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## SUMMARIES

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AUTHORS TITLE	REFERENCE	ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	Locus	AX147742
Goldspink,G.R. and Johnson,I.R. Use of the insulin-like-growth factor i isoform mgf for the treatment of neurological disorders	Eukaryota; Metazoa; Chordata; Crantata; Vertebrata; Eutleteoscom.; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 517)	Homo sapiens	human.		AX147742.1 GI:14346787	AX147742	Sequence 1 from Patent WO0136483.	AX147742 517 bp DNA linear PAT 08	
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Goldspink, G.D. and Terenghi, G.B. Repair of nerve damage Patent: WO 0185781-A 1 15-NOV-2001:
                                                                                                     AX300779 517 bp
Sequence 1 from Patent WO0185781.
AX300779 GI:17382060
                                                          Eukaryota;
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Eukaryota; Metazoa; Chordata; Lepor
Mammalia; Eutheria; Lagomorpha; Lepor
1 (bases 1 to 523)
Coldspink,G.R. and Johnson,I.R.
Use of the insulin-like-growth factor
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DECCFRSCDLRRLEMYCAPLKPAKSARSVRAQRHTDMPKTQKYQPPSTNKNTKSQRRK
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/db_xref="taxon:9606"
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Pred. No. 5e-
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RESULT 4
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AX300783
AX300783.1
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Goldspink, G.D. and Terenghi, G.B. Repair of nerve damage Patent: WO 0185781-A 5 15-NOV-2001;
                                                         Oryctolagus cuniculus
Eukaryota; Metazoa; C
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University
                                             Eukaryota; Metazoa; Chordata; C:
Mammalia; Eutheria; Lagomorpha;
                                                                                                                                           AX300783
                                                                                 rabbit
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)1; Conservative
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/db_xref="GI:14346792"
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/db_xref="taxon:9986"
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                                                         Craniata; Vertebrata;
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AX147754
Sequence 13
AX147754
AX147754.1
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus. 1 (bases 1 to 471)
Goldspink,G.R. and Johnson,I.R.
Use of the insulin-like-growth factor i isoform mgf for treatment of neurological disorders
                                                                                               Oryctolagus cuniculus
Eukaryota; Metazoa; Chordata;
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/db_xref="taxon:9986"
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                                                                                                               AX300791
Sequence
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University College London (GB)
Location/Qualifiers
                                          Oryctolagus cuniculus
Eukaryota; Metazoa; Chordata; Ci
Mammalia; Eutheria; Lagomorpha;
            Goldspink,G.D. and Terenghi,G Repair of nerve damage
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 Patent:
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                                                                                                                                                                                                                                                                                                   GATGTA-GAAGACCCTTCTGAGGAGTGAAGAAGGACAGGCCACCGCAGGACCCTTTGCTC
                                                                                                                                                                                                AGATGGCATTTCCCCCCAATGAAATACACAAGTAAAACAT 469
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of nerve damage
WO 0185781-A 13 15-NOV-2001;
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/db_xxef="GI:1434853"
/taasslatton="GpETL/GAEL/VDALQFVCGDRGEYFNKPTGYGSSSRRAPQTGIV
/translatton="GpETL/GAEL/VDALQFVCGDRGEYFNKPTGYGSSSRRAPQTGIV
DECCERSCDLRRLEMYCAPLKPAKAARSVRAQRHTDMPKTQKEVHLKNTSRGSAGNKN
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/db_xref="taxon:9986"
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Best Local Similarity Matches 455; Conserv
Homo sapiens.

Homo sapiens

Eukaryota, Metazoa, Chordata, Cr.

Bukaryota, Metazoa, Chordata, Cr.

Mammalia, Eutheria, Primates, Cal

1 (bases 1 to 444)

Chew, S. L. Lavender, P., Clark, A.,

An alternatively spliced human in
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                                                                                                                                                   Human alternatively spliced human (IGF-1) mRNA, partial cds. U40870 U40870.1 GI:1100902
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/translation="GPETICGAELVDALQEVRAQRHTDMPKTQKEVHLKNTSRGSAGNKN
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Clark, A.J.

and Ross, R.J

insulin-like

growth factor-I

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356; Conserv
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Chew, S.L.
Direct Submission
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Bartholomew's Hospital Medical College, West Smithfield, London,
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 ATGCCCAAGACCCAGAAGTATCAGCCCCCATCTACCAACAAGAACACGAAGTCTCAGAGA
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/db_xref="taxon:9606"
/clone="pc4"
/tissue_type="liver"
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./gene="IGF-I"
/number=5
395..420
/gene="IGF-I"
/number=6
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/gene="IGF-I"
/note="alternatively spliced; previously, exon 5 and 6
/note="alternatively spliced; this transcript
were thought to be mutually exclusive; this transcript
splices from exon 5 into exon 6; the alternatively spliced
transcript would continue with exon 5 to the polyA signal"
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FVCGDRGFYFNKPTGYGSSSRRAPQTGIVDECCFRSCLLRRLEMYCAPLKPAKSARSV
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/protein_id="AAA96152.1"
/db_xref="GI:1100903"
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/gene="IGF-I"
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Pred. No. 5.6e-103;
0; Mismatches 1;
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Оу Дъ 1	Query Mat Best Loca Matches	exon BASE COUNT ORIGIN	† )	CDS	gene	FEATURES source	JOURNAL MEDLINE PUBMED	TITLE	REFERENCE AUTHORS	PUBMED REFERENCE AUTHORS TITLE JOURNAL		-	20.	Qy 3(
1 GGACCGGAGACGCTCTGCGGGGCTGAGCTGGTGGATGCTCTTCAGTTCGTGTGGAGAC 60	ch 65.6%; Score 344.2; DB 9; Length 616; al Similarity 87.3%; Pred. No. 2.4e-99; 455; Conservative 0; Mismatches 13; Indels 53; Gaps 5;	159	) + 	/gene="IGF-L" 1462 10.3.11011"	ef="tage="tage="type tage="tage="tage="	Location/Qualifiers 1. 616	53 (11), 247	von Holst, H. and Sara, v. Characterization of insulin-like growth factor 1 in human primary brain tumors	STOCKHOLM, SWEDEN 3 (bases 1 to 616) Sandberg-Nordqvist,A.C., Stahlbom,P.A., Reinecke,M., Collins,V.P.,	1372070 (bases 1 to 616) 2 (bases 1 to 616) Sandberg Nordqvist, A.C. Direct Submitssion Submitted (19-NOV-1990) A.C.Sandberg Nordqvist, KAROLINSKA INST'S DEPT OF PATHOLOGY, KAROLINSKA HOSPITAL, BOX 605 00, S-104 01	Sandberg Nordqvist,A.C., Stahlbom,P.A., Lake,M. and Sara,V.K. Characterization of two cDNAs encoding insulin-like growth factor 1 (ICF-1) in the human fetal brain Brain Res. Mol. Brain Res. 12 (1-3), 275-277 (1992) 92186627	oordata; Craniata; Vertebrata; Euteleo imates; Catarrhini; Hominidae; Homo.	HSIGFIA 616 bp mRNA linear PRI 29-NOV-1993 H.sapiens mRNA for IGF-la X56773 S61841 X56773.1 GI:32989	01 AGGAAAGGAAGTACATTTGAAGAACACAAGTAGAGGGAGTGCAGGAAACAAGAACTA 357 

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Prognostic classification of breast cancer
Patent: WO 0210436-A 31 07-FEB-2002;
THE BRIGHAM AND WOMEN'S HOSPITAL, INC. (US
                                                                                                                                                                                                                                                                                                                                                Homo sapiens
Eukaryota; Metazoa;
Mammalia; Eutheria;
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            ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTAT
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ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTAT
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/db_xref="taxon:9606"
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                                                                                                                                                  Score 344.2; DB 6;
Pred. No. 3.4e-99;
0; Mismatches 13;
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Matches 455; Conservative
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Gene expression profiles in liver cancer
Patent: WO 0229103-A 3742 11-APR-2002;
GENE LOGIC INC (US)
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Sequence 3742 from Patent
AX411095
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Mammalia; |
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                                                AGGGGCTTTTATTTCAACAAGCCCACAGGGTATGGCTCCAGCAGTCGGAGGGGCCCTCAG 120
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                       ATGCCCAAGACCCAGAAGTATCAGCCCCCATCTACCAACAAGAACACGAAGTCTCAGAGA
                                                                                                                                                        AGGGGCTTTTATTTCAACAAGCCCACAGGGTATGGCTCCAGCAGTCGGAGGGCGCCTCAG
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  ATGCCCAAGACCCAG-
                                                                                                    ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTAT
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                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
/note="EMBL/GenBBank Accession No.
1415 c 1240 g 2275 t
                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Metazoa;
Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GI:21443800
                                                                                                                                                                                                                                                                                                                                                                            .7260
                                                                                                                                                                                                                                                                       66.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chordata;
Primates;
                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                       Score 344.2; DB 6
Pred. No. 3.4e-99;
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1t W00229103
                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                             Gaps
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KEYWORDS
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MEDLINE
PUBMED
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TITLE
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                                            sig_peptide
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X57025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Direct Submission
Submitted (18-DEC-1990)
CHEMISTRY, UNIVERSITY OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Steenbergh, P.H., Koonen-Reemst, A.M., Sussenbach, J.S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 7260)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               THE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                2018498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Complete nucleotide sequence of the high molecular weight
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Steenbergh, P.H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      insulin-like growth factor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TGCAC-AGTTACCTG-TAAACATTGGAATACCGGCCAAAAAATAAGTTTTGATCACATTTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GATGTAGGAAGACCCTCCTGAGGAGTGAAGAGTGACATGCCACCGCAGGATCCTTTGCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GATGTA-GAAGACCCTTCTGAGGAGTGAAGAAGGACAGGCCACCGCAGGACCCTTTGCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ----AAGGAAGTACATTTGAAGAACGCAAGTAGAAGGAGTGCAGGAAACAAGAACTACAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AGGAAAGGAAGTACATTTGAAGAACACAAGTAGAGGGAGTGCAGGAAACAAGAACTACAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TGCACGAGTTACCTGTTAAACTTTGGAACACCTACCAAAAAATAAGTTTGATAACATTTA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NETHERLANDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mRNA
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                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
/chromosome="12 q22-24.1"
/tissue_type="liver"
/dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Biophys. Res. Commun.
                                                        /translation="MGKISSLPTQLfKCCFCDFLKVKMHTMSSSHLfYLALCLLTFTS
SATAGPETLCGAELVDALQfYCGDRGFYFNKFTGYGSSSRRAPQTGIVDECCFRSCDL
RRLEMYCAPLKPAKSARSVRAQRHTDMPKTQKEVHLKNASRGSAGNKNYRM"
                                                                                                              /evidence=experimental
/product='insulin'like growth factor
/protein_id="CAM40342.1"
/db_xref="GI:33008"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GI:33007
   /evidence=experimental
                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                           167. .310
/gene="IGF-I"
                                                                                                                                                                      /gene="IGF-I"
/codon_start=1
                                                                                                                                                                                                     /evidence=experimental
167. .628
                                                                                                                                                                                                                                /number=1
                                                                                                                                                                                                                                                                                     /gene="IGF-I"
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                                                                                                  /db_xref="SWISS-PROT:P01343"
                                                                                                                                                                                                                                              /gene="IGF-I"
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455; Conser
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                  AAAGAT-GGCATTTCCCCCAATGAAATACACAAGTAAACAT 517
                                                                           TGCAC-AGTTACCTG-TAAACATTGGAATACCGGCCAAAAAATAAGTTTGATCACATTTC 477
                                                                                                                                          GATGTA-GAAGACCCTTCTGAGGAGTGAAGAAGGACAGGCCACCGCAGGACCCTTTGCTC
                                                                                                                                                                                                                                                             ATGCCCAAGACCCAGAAGTATCAGCCCCCATCTACCAACAAGAACACGAAGTCTCAGAGA
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                                                                                                                   GATGTAGGAAGACCCTCCTGAGGAGTGAAGAGTGACATGCCACCGCAGGATCCTTTGCTC
                                                                                                                                                                        ----AAGGAAGTACATTTGAAGAACGCAAGTAGAGGGAGTGCAGGAAACAAGAACTACAG
                                                                                                                                                                                                   AGGAAAGGAAGTACATTTGAAGAACACAAGTAGAGGGAGTGCAGGAAACAAGAACTACAG
    TGCACGAGTTACCTGTTAAACTTTGGAACACCTACCAAAAAATAAGTTTGATAACATTTA
                                                                                                                                                                                                                                  ATGCCCAAGACCCAG-----
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/evidence-experimental
/rpt_family="AluI"
7205 7210
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/number=3
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1415 c 1240 g 2275
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569. .7236
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387. .568
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/evidence=experimental
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/gene="IGF-I"
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861. .865
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="7.6 kb mRNA"
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Pred. No. 3.4e-99;
0; Mismatches 13
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ACCESSION
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                                        AAAGATGGGCGTTTCCCCCAATGAAATACACAAGTAAACAT 538
                                                                                                              TGCAC-AGTTACCTG-TAAACATTGGAATACCGGCCAAAAAATAAGTTTGATCACATTTC
                                                                                                                                                                                                                          ACAGGTATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTAT
                                                                  AAAGAT-GGCATTTCCCCCCAATGAAATACACAAGTAAACAT 517
                                                                                                                                                     GATGTAGGAAGACCCTCCTGAGGAGTGAAGAGTGACATGCCACCGCAGGATCCTTTGCTC
                                                                                                                                                                                                           ----AAGGAAGTACATTTGAAGAACGCAAGTAGAGGGAGTGCAGGAAACAAGAACTACAG
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                                                                                                                                                                                                                                                                                                                          TGCGCACCCTCAAGCCTGCCAAGTCAGCTCGCCTGTCCGTGCCCAGCGCCACACCGAC
                                                                                               TGCACGAGTTACCTGTTAAACTTTGGAACACCTACCAAAAAATAAGTTTGATAACATTTA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /product="IGF-1 precursor"
/protein_id="CaAO1955.1"
/protein_id="CaAO1955.1"
/db_xref="Gui+$12993"
/translation="MALCLLIFTSSATAGPETLCGAELVDALQFVCGDRGFYFNKPTG
/translation="MALCLLIFTSSATAGPETLCGAELVDALQFVCGDRGFYFNKPTG
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YGSSSRRAPOTGIVDECCFRSCDLRRLEMYCAPLKPAKSARSVRAQRHTDMPKTQKEV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /product="IGF-1"
167 c 181
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              145
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.9e-99;
les 14;
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patent GB2241703.
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MEDLINE
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Matches 454
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Jansen,M., van Schaik,F.M., Ricker,A.T., Bullock,B., Gabbay,K.H., Nussbaum,A.L., Sussenbach,J.S. and Van Sequence of cDNA encoding human insulin-like growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          X00173.1 GI:33015
growth factor; insulin super
signal peptide; somatomedin.
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Location/Qualifiers
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ilarity 87.18;
Conservative
                                                                                    - AAGGAAGTACATTTGAAGAACGCAAGTAGAGGGAGTGCAGGAAACAAGAACTACAG
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/codon_start=1
/product="insulin-like growth factor lA precursor"
/protein_id="caa24998.1"
/protein_id="caa24998.1"
/db_xref="gi:33016"
/db_xref="swiss-prot-gol343"
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174 c
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12. .473
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Primates;
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Pred. No. 8e-9
0; Mismatches
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Catarrhini; Hominidae
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n den Brande,J.L.
h factor i
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3683205
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Human insulin-like growth factor I and II messenger RNA: isolation
of complementary DNA and analysis of expression
Meth. Enzymol. 146, 239-248 (1987)
                                                                                                                                                                                                                                                                                        Chromosome 12q23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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Human (adult) liver, cDNA to mRNA.
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                                                                           ACAGGCATCGTGGATGAGTGCTGCTGCGGAGCTGTGATCTAAGGAGGCTGGAGATGTAT
                                                                                                                               AGGGGCTTTTATTTCAACAAGCCCACAGGGTATGGCTCCAGCAGTCGGAGGGGCGCCTCAG 120
                                                                                                                                                                       GGACCGGAGACGCTCTGCGGGGCTGAGCTGGTGGATGCTCTTCAGTTCGTGTGTGGAGAC
                                                                                                                                                                                                   GGACCGGAGACGCTCTGCGGGGCTGAGCTGGTGGATGCTCTTCAGTTCGTGTGGAGAC 60
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             AGGGGCTTTTATTTCAACAAGCCCACAGGGTATGGCTCCAGCAGTCGGAGGGCGCCTCAG
TGCGCACCCCTCAAGCCTGCCAAGTCAGCTCGCTCTGTCCGTGCCCCAGCGCCACACCGAC
                                                        ACAGGTATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTAT
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                                                                                                                                                                                                                                                                                                     193
                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                  /gene="IGF1"
/note="insulin-like growth factor I signal
156. .365
/gene="IGF1"
/product="insulin-like growth factor I"
174 c 183 g 178 t
                                                                                                                                                                                                                                                                                                                                                                                                        /translation="MHTMSSSHLFYLALCLLTFTSSATAGPETLCGAELVDALQFVCG
DRGFYFNKPTGYGSSSRRAPQTGIVDECCFRSCDLRRLEMYCAPLKPAKSARSVRAQR
HTDMPKTQKEVHLKNASRGSAGNKNYRM"
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 /db_xref="GDB:G00-120-081"
                                                                                                                                                                                                                                                                                                                                                                                                                                                              /protein_id="AAA52543.1"
/db_xref="GI:183120"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /codon_start=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note="insulin-like growth
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Primates;
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Pred. No. 8e-99;
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RESULT 1
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M27544.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 1076)
Le Bouc,Y., Dreyer,D., Jaeger,F., Binoux,M. and Sondermeye:
Complete characterization of the human IGF-I nucleotide sec
isolated from a newly constructed adult liver cDNA library
FEBS Lett. 196 (1), 108-112 (1986)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HUMIGFI 1076 bp mRN. Human insulin-like growth factor mRNA,
                            /product="insulin-like 283 a 251 c 239 g 3 Chromosome 7p13-p12.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                              293.
                                                                                                                                              149.
                                                                                                                                                          /db_xref="GDB:G00-120-075"
/translation="MGKISSLPTQLFKQCFCDFLKVKMHTMSSSHLFYLALCLLTFTS
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RRLEMYCAPLKPAKSARSVRAQRHTDMPKTQKEVHLKNASRGSAGNKNYRM"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                         /codon_start=1
/protein_id="AAA52787.1"
/db_xref="GI:306927"
                                                                                                                                                                                                                                                                                                        149.
                                                                                                                                                                                                                                                                                                                                                                  /gene="IGFBP1"
/note="IGF mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                 /gene="IGFBP1"
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                                                                                                            /note="insulin-like
                                                                                                                           /gene="IGFBP1"
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                                                                               /gene="IGFBP1"
                                                                                                                                                                                                                                                                      'note="insulin-like growth factor precursor"
                                                                                                                                                                                                                                                                                       'gene≖'
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numan IGF-I nucleotide sequer
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Listing first 45 summaries
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                                                                                          Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.
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/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1982.DAT:*
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7423.602 Million cell updates/sec
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Length D		ID		Description
517	22	AAD06398		Human IGF-I isofor
	24	AAS16877		Human mechano-grow
•	22	AAD06400		Rabbit IGF-I isofo
523	24	AAS16879		Rabbit mechano-gro
	18	AAT84893		Rabbit insulin lik
	22	AAD06405		Rabbit liver-type
	24	AAS16884		Rabbit insulin-lik
818	æ	AAN70436		Sequence encoding
_		ABK84583		Human cDNA differe
1000 1000 1000 1000 1000 1000 1000 100	Match Length D 100.0 517 100.0 517 90.4 523 90.4 553 90.4 553 90.4 471 73.0 471 66.6 818 66.6 7260	Length DB 517 22 517 24 523 24 553 18 553 18 471 24 7260 24	Length DB ID  517 22 AAD06398 517 24 AAS16877 518 24 AAS16809 523 24 AAS16809 523 24 AAS16809 553 18 AAT84893 471 22 AAS168405 471 24 AAS16884 818 8 AAN70436 7260 24 ABK84583	AANO6398 AAS16877 AANJ6400 AAS16879 AATB4893 AATB4893 AANJ6884 AANJ6405 AANJ6405 AANJ6405 AANJ6405 AANJ6405 AANJ6405 AANJ6405 AANJ6405 AANJ6405

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ingle nucleo	AAC7201	ي د	Δ	л	S	Л
eq ID 7 u	AAZ4427	21	N	7	92.	44
q ID 1 use	AAZ4426	21	1	7	92.	ω.
iller toxin	AAT6352	18	Ø	9	02.	42
ynthetic hum	AAQ2330	р ( s )	4	9.	02.	41
insu	AAD3201	24	-	9.	03.	40
alpha	AAD3201	24	S	9.	04.	39
inant b	AAV2628	19	S	0	208.4	38
1/IGF-	AAQ1356	7	w	0	08.	37
al/IGF	AAQ1356	1	w	0	$\circ$	36
ature	ABA0314	24	$\vdash$	0	$\vdash$	5
Rat insulin-li	AAS1688	24	$\alpha$	7.	47.	34
iver-typ	AAD0640	22	$\alpha$	7	47.	33
iver form	AAX2749	20	UN	8	52	32
poly	ABL9169	24	on	0	58.	31
Human IGF-I cod	AAV4079	19	g	0	58.	.0
H -	AAV5042	19	ത	0	58.	29
ver-	AAD0640	22	-	0	58.	28
ω	AASl	24	317	50.0	258.4	27
encod	AAQ4780	14	7		71.	26
an cDN	AAS1469	22	س	5	85.	5
smid pIGO10	AAX8805	20	34	ū	86.	24
lasmid pIG033	AAX8805	20	70		86.	23
xpected se	AAV4079	19	9		86.	22
lasmid pIG05	AAV5042	19	60	çn	86.	21
ctual se	AAV4079	19	9	ÇI	86	20
lasmid pIG05	AAV5042	19	59		86.	9
e encod	AAN7043	œ	13	ω.	08	8
t mechano-g	AAS1687	24	ω		25.	7
IGF-	AAD063	22	ω	ν.	25.	9
repro-s	AAN6049	7	$\sim$	5	39.	5
າ insuli	AAT8489	18	$\sim$	5	42.	4
į	ABK3556	24	26		44.	Lω
ndome	ABK355	24	O)	<u>ა</u>	44.	12
man benign	ABK6481	24	26		44.	Ε
Gene #3742 use	ABN9724	24	26	<u>ი</u>	44	0

## ALIGNMENTS

10-AUG-2001 AAD06398;

(first entry)

AAD06398 standard; cDNA; 517 BP

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ARDOGSS 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human IGF-I isoform mechano-growth factor (MGF) cDNA
                                                          25-MAY-2001
                                                                                                                                                                  WO200136483-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                         /*tag= a
/product= "Mechano-growth factor (MGF)"
/note= "This region comprises exons 3-6.
not include start codon"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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Query Match
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Matches 517
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            toxin, motoneurone trauma, a motoneurone lesion or nerve damage, an injury that affects motoneurones, motoneurone loss associated with aging, autosomal or sex-linked muscular dystrophy, diabetic neuropathy, peripheral neuropathies, Alzheimer's disease and Parkinson's disease. The present sequence is human IGF-I isoform MGF DNA. MGF is a muscle isoform having extracellular (Ec) domain, hence also referred as IGF-I-Ec. The MGF protein comprises amino acid sequences encoded by nucleic acid sequence of IGF-I exons 4, 5 and 6 in the reading frame
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; Page 49-50; 66pp; English
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                                                                           GATGTAGAAGACCCTTCTGAGGAGTGAAGAAGGACAGGCCACCGCAGGACCCTTTGCTCT
                                                                                                                                                      GATGTAGAAGACCCTTCTGAGGAGTGAAGAAGGACAGGCCACCGCAGGACCCTTTGCTCT
                                                                                                                                                                                                                                                                                AGGAAAGGAAGTACATTTGAAGAACACAAGTAGAGGGAGTGCAGGAAACAAGAACTACAG
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GCACAGTTACCTGTAAACATTGGAATACCGGCCAAAAAATAAGTTTGATCACATTTCAAA
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llarity 100.0%;
Conservative (
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Pred. No. 2e-146;
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The invention relates to the use of an insulin-like growth factor I (IGF-I) isoform, known as mechano-growth factor (WGF), in the manufactor a medicament for treating nerve damage in the peripheral nervous system, or for treating nerve damage by localising MGF at the site of damage. The nerve damage may include severing of a nerve. The treatment may be combined with another treatment (such as a polypeptide growth
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                                                                                                                                                                                                                                                      Use of insulin-like growth factor I (IGF-I) isoform known as mechano-growth factor which is encoded by IGF-I exons 4,5,6 and ability to reduce motoneuron loss in response to nerve avulsion,
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                                                                                     """ insulin-like Growth Factor-I; MGF;
""" amyotrophic lateral sclerosis; spinal muscular atrophy; mulcular atrophy; poliomyelitis; post-polio syndrome; toxin; motoneurone disorder; nerve damage; autosomal muscular dystrophy; diabetic neuropathy; sex-linked muscular dystrophy; peripheral neuropathy; Alzheimer's disease; Parkinson's disease; sa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            factor other than MGF) that prevents or diminishes degeneration of the target organ (for example, muscle) which the damaged nerve innervates, whereby the treatment of the muscle with MGF or a polynuclectide encoding MGF prevents or diminishes degeneration. The method is useful for treating neurological disorders, preferably motorneuron disorders. These methods can reduce motoneuron loss by 20% or greater in response to nerve avulsion. This sequence represents cDNA encoding the human MGF.
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                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to use of mechano-growth factor (MGF), an Insulin-like Growth Factor-I (IGF-I) isoform in the manufacture of a medicament for the treatment of neurological disorder. The MGF is capable of reducing motoneurone loss by 20% or greater in response to nerve avalsion, and effects motoneurone rescue, preferably adult motoneurone rescue. The MGF polypuclectide and polypeptide are useful in the including a disorder of motoneurones and/or neurodegenerative disorder, including a disorder of motoneurones and/or neurodegenerative disorder, e.g., amyotrophic lateral sclerosis, spinal muscular atrophy, progressive spinal muscular atrophy, infantile or juvenile muscular atrophy, progressive poliomyelitis or post-polio syndrome, a disorder caused by exposure to a toxin, motoneurone trauma, a motoneurone lesion or nerve damage, an injury that affects motoneurones, motoneurone loss associated with aging, autosomal or sex-linked muscular dystrophy, diabetic neuropathy. The present sequence is rabbit IGF-I sicoform MGF coMA. MGF is a muscle isoform having extracellular (Ec) domain, hence also referred as IGF-I-Ec. The MGF protein comprises amino acid sequences encoded by complete acid sequence of IGF-I exons 4, 5 and 6 in the reading frame
                                                                                                                                                                                                                                                                                                                  Query Match
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Matches 501
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                                                                              AGGGGCTTTTATTTCAACAAGCCCACAGGATACGGCTCCAGCAGTCGGAGGGCACCTCAG
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                          ATGCCCAAGACCCAGAAGTATCAGCCCCCATCTACCAACAAGAACACGAAGTCTCA---G
                                                                                                                        ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTGAGGAGGCTGGAGATGTAC
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/product= "Mechano-growth factor (MGF)"
/note= "This region comprises exons 3-6.
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Use of insulin-like growth factor I (IGF-I) isoform known as mechano-growth factor which is encoded by IGF-I exons 4.5.\% and ability to reduce motoneuron loss in response to nerve avulsion,
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Best Local S
Matches 501
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (IGF I) Isoform, known as mechano-growth rather the peripheral nervous of a medicament for treating nerve damage in the peripheral nervous system, or for treating nerve damage by localising MGF at the site of damage. The nerve damage may include severing of a nerve. The treatment may be combined with another treatment (such as a polypeptide growth factor other than MGF) that prevents or diminishes degeneration of the target organ (for example, muscle) which the damaged nerve innervates, whereby the treatment of the muscle with MGF or a polynucleotide encoding MGF prevents or diminishes degeneration. The method is useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MGF prevents or diminishes degeneration. The method is useful for treating neurological disorders, preferably motorneuron disorders. These methods can reduce motoneuron loss by 20% or greater in response to nerve avulsion. This sequence represents cDNA encoding the rabbit MGF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to the use of an insulin-like growth (IGF-I) isoform, known as mechano-growth factor (MGF), in t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                treat nerve damage
                                                 Rabbit insulin like growth factor 1 encoding cDNA
                                                                                                                                                 AAT84893 standard; cDNA;
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                                                                                                                                                                                                                                                   CAAAGATGGCATTTCCCCCCAATGAAATACACAAGTAAACAT
                                                                                                                                                                                                                                                                                                                                                                                       CAGGATGTA-GAAGACCCTTCTGAGGAGTGAAGAAGGACAGGCCACCGCAGGACCCTTTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TGTGCACCCCTCAAGCCGGCAAAGGCAGCCCGCTCCGTGCCCAGCGCCCACACCGAC
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                                                                                                                                                                                                                                     CAAAGATGGCATTTCCCCCCAATGAAATACACAAGTAAACAT
                                                                                                                                                                                                                                                                                                                                                                     CAGGATGTAGGAAGACCCTTCTGAGGAGTGAAGAAGGACAGGCCACCGCAGGACCCTTTG
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                                                                                                                                                                                                                                                                                                   CTCTGCACAGTTACCTGTAAACATTGGAATACCGGCCAAAAAATAAGTTTGATCACATTT
                                                                                                                                                                                                                                                                                                                                   CTCTGCACAGTTACCTGTAAACATTGGAATACCGGCCAAAAAATAAGTTTGATCACATTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AGAAGGAAAGGAAGTACATTTGAAGAACACAAGTAGAGGGAGTGCAGGAAACAAGAACTA
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                                                                                   (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 467.4; DB 24;
Pred. No. 2.1e-131;
0; Mismatches 16;
                                                                                                                                                 ВP
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Insulin like growth factor 1; IGF-1; Ec peptide; muscle disorder; heart; neuromuscular disease; primer; ss.

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                                                                                                                                                                                                                                                                                                                                                    Matches 501;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   A use of insulin like growth factor I (IGF-1) has been developed, and is characterised by the presence of the Ec peptide, or a functional equivalent, in the treatment or therapy of a human or animal The IGF-1 polypeptide can be used to treat muscular disorders, e.g. Duchenne or Becker muscular dystrophy, autosomal dystrophies and related progressive skeletal muscle weakness and wasting, muscle atrophy in ageing humans, spinal cord injury induced muscle atrophy and neuromuscular diseases, and cardiac disorders, e.g. diseases where promotion of cardiac muscle protein synthesis is a beneficial treatment, cardiomyopathies and acute heart failure or insult, specifically myocardiits or myocardial infarction. It can also be used to promote bone fracture healing and maintenance of bone in old age. The present sequence encodes rabbit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Use of insulin like growth factor I characterised by presence of Ec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Goldspink
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                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 553 BP; 159 A; 142 C; 147 G; 105 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (UNLO ) ROYAL
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DB; AAW23301.
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                                                                                                                                                                                                                                                                                                                                                                                                                                      used in the present specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          conditions or neuromuscular diseases
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                                                                                                                                                                                                                                                                                                                      GGACCGGAGACGCTCTGCGGGGCCTGAGCTGGTGGATGCTCTTCAGTTCGTGTGGAGAC
                                                                                                                                                                                                                                                                                            GGACCGGAGACGCTCTGCGGTGCTGAGCTGGTGGATGCTCTTCAGTTCGTGTGGAGAC
                                                                            ATGCCCAAGACCCAGAAGTATCAGCCCCCATCTACCAACAAGAACAACAAGTCTCA---
                                                                                                                                     TGCGCACCCCTCAAGCCTGCCAAGTCAGCTCGCTCTGTCCGTGCCCAGCGCCCACACCGAC
                                                                                                                                                                         ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTGAGGAGGCTGGAGATGTAC
                                                                                                                                                                                          ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTAT
                                                                                                                                                                                                                                  AGGGGCTTTTATTTCAACAAGCCCACAGGATACGGCTCCAGCAGTCGGAGGGCACCTCAG
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AGAAGGAAAGGAAGTACATTTGAAGAACACAAGTAGAGGGAGTGCAGGAAACAAGAACTA
                                                                                                                 AGAAGGAAAGGAAGTACATTTGAAGAACACAAGTAGAGGGAGTGCAGGAAACAAGAACTA
                                                         ATGCCCAAGACTCAGAAGTATCAGCCTCCATCTACCAACAAGAAAATGAAGTCTCAGAGG
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                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                90.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               English.
                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                Score 467.4; DB 18 Pred. No. 2.1e-131;
                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                 DB 18;
                                                                                                                                                                                                                                                                                                                                                    16;
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                                                                                                                                                                                                                                                                                                                                                                              Length 553;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rabbit; IGF-I isoform; Insulin-like Growth Factor-I; MGF; mechano-growth factor; neurological disorder; neurodegenerative disorder; amyotrophic lateral sclerosis; spinal muscular atrophy; muscular atrophy; poliomyelitis; post-polio syndrome; toxin; motoneurone disorder; nerve damage; autosomal muscular dystrophy; diabetic neuropathy; sex-linked muscular dystrophy; peripheral neuropathy; sex-linked fuscular dystrophy; peripheral neuropathy; ss. Alzheimer's disease; Parkinson's disease; liver; L.IGF-I; ss.
The present invention relates to use of mechano-growth factor (MGF), an Insulin-like Growth Factor-I (IGF-I) isoform in the manufacture of a medicament for the treatment of neurological disorder. The MGF is capable of reducing motoneurone loss by 20% or greater in response to nerve avulsion, and effects motoneurone rescue, preferably adult motoneurone rescue. The MGF polynucleotide and polypeptide are useful in the manufacture of a medicament for the treatment of a neurological disorder...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oryctolagus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rabbit liver-type IGF-I isoform (L.IGF-I) cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-AUG-2001
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                                                                                                                                                                                                                                                                                                                           Use of mechano-growth factor, an isoform of Insulin-like Growth Factor-I, capable of reducing motoneurone loss, in the manufacture medicament for the treatment of neurological disorder \dot{}
                                                                                                                                                                                                                                                                                                                                                                                                                                                               P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAY-2001
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                                                                                                                                                                                                                                                                    Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (UNLO ) UNIV COLLEGE LONDON
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DB; AAE02452, AAE02456
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                                                                                                                                                                                                                                                                Page 59-60; 66pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000WO-GB04354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Johnson
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/transl_except= (pos:25..27, aa:Gln)
/note= "These translation exceptions occur while decoding the alternative version of the protein (AAE02456).
The CDS comprises exons 3, 4 and 6 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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Matches 455
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        toxin, motoneurone trauma, a motoneurone lesion or nerve damage, an injury that affects motoneurones, motoneurone loss associated with ag autosomal or sex-linked muscular dystrophy, diabetic neuropathy, peripheral neuropathies, Alzheimer's disease and Parkinson's disease. The present sequence is rabbit liver type IGF-I isoform (L.IGF-I) cDN The L.IGF-I protein comprises amino acid sequences encoded by nucleic acid sequence of IGF-I exons 4 and 6.
                                                                 Rabbit; mechano-growth factor; insulin-like growth factor I; IGF-I; MGF; neuroprotective; nerve damage; peripheral nervous system; nerve severing; muscle; neurological disorder; motoneuron loss; motorneuron disorder; sorter; ss; nerve avulsion; insulin-like growth factor I liver-type isoform; L.IGF-I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       including a disorder of motoneurones and/or neurodegenerative disorder, e.g., amyotrophic lateral sclerosis, spinal muscular atrophy, progressi spinal muscular atrophy, infantile or juvenile muscular atrophy poliomyelitis or post-polio syndrome, a disorder caused by exposure to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 471
                                    Oryctolagus
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                                                                                                                                                 Rabbit insulin-like growth factor I liver-type isoform (L.IGF-I)
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   Location/Qualifiers
                                                                                                                                                                                                                                                  cDNA; 471
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Pred. No. 4.1e-104;
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Best Local S
Matches 455
                                                                                                                                                                                                                                                                              The invention relates to the use of an insulin-like growth factor I (IGF-I) isoform, known as mechano-growth factor (MGF), in the manufacture of a medicament for treating nerve damage in the peripheral nervous system, or for treating nerve damage by localising MGF at the site of damage. The nerve damage may include severing of a nerve. The treatment may be combined with another treatment (such as a polypeptide growth factor other than MGF) that prevents or diminishes degeneration of the target organ (for example, muscle) which the damaged nerve innervates, whereby the treatment of the muscle with MGF or a polynucleotide encoding MGF prevents or diminishes degeneration. The method is useful for treatment of the muscle with MGF or a polynucleotide encoding MGF prevents or diminishes degeneration. The method is useful for treating neurological disorders, preferably mortonneuron disorders. These methods can reduce motoneuron loss by 20% or greater in response to nerve avulsion. This sequence represents cDNA encoding the rabbit insulin-like growth factor I liver-type isoform (L.IGF-I) used in experiments on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          exon
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Use of insulin-like growth factor I (IGF-I) isoform known mechano-growth factor which is encoded by IGF-I exons 4.5 ability to reduce motoneuron loss in response to nerve av
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                                                                                                                                                  ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTAT
                                                                                             AGGGGCTTTTATTTCAACAAGCCCACAGGGTATGGCTCCAGCAGTCGGAGGGCGCCTCAG
                                                                                                                                                                                                                                               471
TGCGCACCCCTCAAGCCTGCCAAGTCAGCTCGCTCTGTCCGTGCCCAGCGCCCACACCGAC
                           ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTGAGGAGGCTGGAGATGTAC
                                                                               AGGGGCTTTTATTTCAACAAGCCCCACAGGATACGGCTCCAGCAGTCGGAGGGCACCTCAG
                                                                                                                                    GGACCGGAGACGCTCTGCGGTGCTGAGCTGGTGGATGCTCTTCAGTTCGTGTGTGGAGAC
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GRINSTEAD MEDICAL
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                                                                                                                                                                                      Score 377.2; I
Pred. No. 4.1e
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                                                                       A 42 base oligonucleotide corresponding to the DNA sequence encoding amino acids 10 to 23 of mature human IGF-I was synthesized (AAN70437). The radiolabeled 42 mer was then employed to screen for IGF-I containing DNA sequences in a human liver cDNA library. Insulinlike growth factors-1A and -1B cDNAs were isolated from a human cDNA library by using lambdagt II (AAN70435, AAN70436). The human IGF-I genomic gene was isolated and mapped. It encodes at least two preproinsulin-like growth factor-1 proteins. An essentially pure proproinsulin-like growth factor-1 protein comprising the sequence
                                                                                                                                                                                                                                                   New pre-pro-insulin-like growth factor-1 protein - obtd. recombinant DNA procedures for use as growth promoters f enhancing lactation, for stimulating cell proliferation
                               Sequence 818
                                                                                                                                                                                                                          Example; Fig 5; 59pp; English.
                                                                                                                                                                                                                                                                                                                WPI; 1987-200203/29
                                                                                                                                                                                                                                                                                                                                                                         (UNIW ) UNIV OF WASHINGTON
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07-JAN-1986;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Growth promoter;
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                                                            acids shown
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86US-0816662
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                               232 A; 186 C; 187 G; 213 T; 0 other;
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                                                           in Figure
   66.6%;
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                                                         or-1 proteins. An essentially pure or-1 protein comprising the sequence six is claimed (AAP70277).
   344.2;
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              Beazer-Barclay Y,
                                                                                                                                                                                                                                                rheumatoid arthritis; glomerulonephritis; asthma; thrombosis; cardiac reperfusion injury; renal reperfusion injury; ARDS; adult respiratory distress syndrome; inflammatory bowel disease;
                                                                                                                                                                                                                                                                                                        Human; ss; granulocytic cell; DNA chip; bacterial infection; viral infection; parasitic infection; protozoal infection;
                                                                       03-OCT-2000; 2000US-237189P
                                                                                                       03-OCT-2001; 2001WO-US30821
                                                                                                                                                               WO200228999-A2
                                                                                                                                                                                                                      Crohn's disease; ulceration;
                                                                                                                                                                                                                                                                                                                                                       Human cDNA differentially expressed in granulocytic cells #1154
                                                                                                                                                                                             Homo
                                                                                                                                                                                                                                                                                            viral infection; parasitic infection; protozoal infection; fungal infection; sterile inflammatory disease; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                       14-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                ABK84583 standard;
                                           (GENE-) GENE LOGIC
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                                                                                                                                                                                                                       ulcerative colitis; periodontal disease; vation; chronic inflammation; allergy.
               Weissman
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detecting granulocyte activation by detecting differential expression of genes associated with granulocyte activation, which serves as diagnostic markers that is useful for monitoring disease states and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 7260 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to detecting (M1) granulocyte (GC) activ. (GCA), by detecting the level of expression of gene(s) (Gs) id DNA chip analysis as given in the specification, and comparing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; SEQ ID No 1154; 114pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
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                                                                                                                                                  TGCGCACCCCTCAAGCCTGCCAAGTCAGCTCGCTCTGTCCGTGCCCAGCGCCACACCCGAC
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AGGAAAGGAAGTACATTTGAAGAACACAAGTAGAGGGAGTGCAGGAAACAAGAACTACAG
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87.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Score 344.2; 1; Pred. No. 1.2e. 0; Mismatches
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BP; 2330 A; 1415 C; 1240 G;

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                         The invention relates to a novel method for diagnosing and detecting the progression of liver cancer, hepatocellular carcinoma or metastatic liver tumour in a patient, and differentiating metastatic liver cancer from the patocellular carcinoma in a patient, involving detecting the level of expression of two or more genes represented in ABN93503-ABN97455 in a ctissue sample. The method of the invention has hepatotropic, and ctissue sample the method is useful for diagnosing and detecting the progression of liver cancer, hepatocellular carcinoma and metastatic liver carcinoma in a patient. The method is useful for identifying expression profiles which serve as useful diagnostic markers as well as markers that can be used to monitor disease states, disease progression, contest the sequence data for this patient did not form part of the printed specification, but was obtained in electronic format directly from WIPO at the proposition of the proposition of the printed specification, but was obtained in electronic format directly from WIPO at the proposition of the printed specification.
                                                                                                                                                                                                                                                                                                                                                                                     Diagnosing and detecting the progression of liver cancer, hepatocellular carcinoma or metastatic liver tumor in a patient, involves detecting the level of expression of two or more genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gene; liver cancer; ds; hepatocellular carcinoma; hepatotropic; metastatic liver tumour; cytostatic; expression profile; disease si disease progression; drug toxicity; drug efficacy; drug metabolism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gene #3742 used
                                                                                                                                                                                                                                                                                                                                    Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-OCT-2001; 2001WO-US30589
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GENE-) GENE LOGIC INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-OCT-2000; 2000US-237054P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABN97244 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 3742;
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Identifying drugs for and diagnosing benign prostatic hyperplasia, by detecting expression levels of one or more genes in prostate cells from patient that are differentially regulated compared to normal prostate cells
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05-JUN-2001;
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N TOBACCO INC
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RESULT 1: ABK35504 ID ABK

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               from the subject that are differentially regulated compared to normal prostate cells. (II) comprises preparing a first gene expression profile of BPH cells or BPH-like cell population, exposing the cells to the agent, preparing a second gene expression profile of the agent exposed cells, and comparing the first and second gene expression profiles. (II) is useful for diagnosing the onset or progression of BPH. (II) is useful for identifying an agent that modulates the onset or progression of BPH. The methods are useful to present information identifying the expression of genes given in the specification in the tissue or cells to the expression of gene in the database, and displaying the expression level of expression level in a tissue or cells to the compared to the expression level in BPH. Agents using (II) are useful for treating BPH or prostate cancer. ABK64105-ABK64860 represent human tax
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a method of diagnosing (1) the onset or progression of benign prostatic hyperplasta (BPH), or screening (1) or identifying an agent that modulates the onset or progression of The method is based on changes in gene expression in BPH tissue isc from patients exhibiting different clinical states of prostate hyperplasta as compared to normal prostate tissue (I) comprises detecting the expression levels of one or more genes in prostate ce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 391-393; 444pp; English
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                     GATGTA-GAAGACCCTTCTGAGGAGTGAAGAAGGACAGGCCACCGCAGGACCCTTTTGCTC
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                                                                                            AGGAAAGGAAGTACATTTGAAGAACAAGTAGAGGGAGTGCAGGAAACAAGAACTACAG
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                                                                      ----AAGGAAGTACATTTGAAGAACGCAAGTAGAGGGAGTGCAGGAAACAAGAACTACAG
                                                                                                                                                                                  ATGCCCAAGACCCAGAAGTATCAGCCCCCATCTACCAACAAGAACACGAAGTCTCAGAGA
                                                                                                                                                                                                                     TGCGCACCCCTCAAGCCTGCCAAGTCAGCTCGCTCTGTCCGTGCCCAGCGCCACACCGAC
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Pred. No. 1.2e-93
0; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to diagnosing endometrial cancer in a subject consumption of a set of nucleic acid molecules or expression of a set of nucleic acid molecules or expression products in an endometrial sample suspected of being cancerous, where the set of nucleic acid molecules comprises at least 2 nucleic acid set of the 50 nucleic acids are used as an array of at least 2 of the 50 nucleic acids bound to a solid substrate. Also included is a solid phase comprising at least 2 antibodies or its antigen binding fragments, that specifically bind at least 2 different polypeptides from the 50 fully defined sequences as given in the specification, fixed to a solid substrate. The methods and arrays are cuseful for the diagnosis of endometrial cancer, selecting and monitoring treatment of endometrial cancer. The present sequence is one of 50 treatment of endometrial cancer. The present sequence is one of 50 treatment of endometrial cancer.
                                                                                                                                                                                                                                                                                                                          Best
                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Diagnosing endometrial cancer comprises determining expression of nucleic acid molecules or expression products that are differentially expressed in normal and malignant endometrium -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; ds; gene; encomet DNA microarray; protein
                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
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                                                                                                                                                                                                                                                                                                                          Local
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                                                                                                                                                                                                                                                                                                                                                                                                                            differentially expressed between cancerous and non-cancerous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Page 85-89; 233pp;
                                                                                                                                                                                                      61
                                                                                                                                                                                                                                                                      1 GGACCGGAGACGCTCTGCGGGGCCTGAGCTGGTGGATGCTCTTCAGTTCGTGTGGAGAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAU84284
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 ATGCCCAAGACCCAGAAGTATCAGCCCCCATCTACCAACAAGAACACGAAGTCTCAGAGA
                                  TGCGCACCCTCAAGCCTGCCAAGTCAGCTCGCTCTGTCCGTGCCCAGCGCCACACCGAC
                                                                 TGCGCACCCCTCAAGCCTGCCAAGTCAGCTCGCTCTGTCCGTGCCCAGCGCGCCACACCGAC
                                                                                                                                                                    AGGGGCTTTTATTTCAACAAGCCCACAGGGTATGGCTCCAGCAGTCGGAGGGCGCCTCAG
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protein microarray.
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Pred. No. 1.2e
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RESULT 13
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                                                                                                                                                                                      Diagnosing breast cancer comprises determining expression of nucleic acid molecules or expression products that are differentially expressed in normal and malignant tissue \cdot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABK35561 standard; DNA;
                                                                                                                                                                     Claim 1;
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                                                                                                                                                                                                                                                            Baak J,
                                                                                                                                                                                                                                                                                                              28-JUL-2000; 2000US-222093P
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MAI; mitotic activity index; cytostatic;
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                                                                                                                                                                                                                                                                                 BAAK
                                                                                                                                                                                                                                                                                          BRIGHAM & WOMENS HOSPITAL INC
                                                                                                                                                                   Page 74-78; 219pp; English
                                                                                                                                                                                                                                                            Mutter GL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AGGAAAGGAAGTACATTTGAAGAACACAAGTAGAGGGAGTGCAGGAAACAAGAACTACAG
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The present invention relates to a method for diagnosing breast cancer in a subject suspected of having endometrial cancer. The method comprises determining the expression of a set of human genes or expression products in an endometrial sample suspected of being cancerous. The human genes of the invention are differentially expressed in breast tumours characterised as high or low MAI without expressed in breast tumours characterised as high or low MAI without expressed in breast tumours. The invention also provides DNA and protein microarrays for analysing the expression of the human genes and their protein products. The methods and arrays are useful for the diagnosis and prognosis of endometrial cancer, selecting and monitoring treatment regimes, and identification of compounds useful for the treatment of endometrial cancer. ABX3551-ABX3581 represent the human genes of the invention that are differentially expressed in breast

cancer tissue

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Matches
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                                              11-MAR-1997;
                                                                                                                                                                                      Human insulin like growth factor 1 Ea isoform encoding cDNA
                                                                                                                                                                                                         14-APR-1998
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                            11-MAR-1996;
                                                               18-SEP-1997.
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                            96GB-0005124
                                             97WO-GB00658
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/product=
                                                                                                                 Location/Qualifiers 26..496
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Pred. No. 1.:
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Query Match
Best Local Simi
Matches 454;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A use of insulin like growth factor I (IGF-1) has been developed, and is characterised by the presence of the Ec peptide, or a functional equivalent, in the treatment or therapy of a human or animal. The IGF-1 polypeptide can be used to treat muscular disorders, e.g. Duchenne or Becker muscular dystrophy, autosomal dystrophies and related progressive skeletal muscle weakness and wasting, muscle atrophy in ageing humans, spinal cord injury induced muscle atrophy and neuromuscular diseases, and cardiac disorders, e.g. diseases where promotion of cardiac muscle protein synthesis is a beneficial treatment, cardiomyopathies and acute infarction. It can also be used to promote bone fracture healing and maintenance of bone in old age. The present sequence encodes human IGF-1 Ea isoform used in the present specification.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 777 BP; 201 A; 193 C; 204 G; 179 T; 0 other;
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                                                           AAAGAT-GGCATTTCCCCCAATGAAATACACAAGTAAACAT 517
                                                                                                                                                       TGCAC-AGTTACCTG-TAAACATTGGAATACCGGCCAAAAAATAAGTTTGATCACATTTC
                                                                                                                                                                                                                                               GATGTAGGAAGACCCTCCTGAGGAGTGAAGAGTGACATGCCACCGCAGGATCCTTTGCTC
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AAAGATGGGCGTTTCCCCCCAATGAAATACACAAGTAAACAT
                                                                                                                         TGCACGAGTTACCTGTTAAACTTTGGAACACCTACCAAAAAATAAGTTTGATAACATTTA
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87.1%;
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Pred. No. 1.
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Query Match
Best Local Similarity
Matches 363; Conserv
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See also AAN60489, AAN60491

(Updated on 03-OCT-2002 to add missing OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 522 BP; 171 A; 147 C; 161 G; 143 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pre:pro insulin-like growth factors I and II - obtd. from the human genome by e.g. screening a cDNA library obtd. from human liver cells.
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P-PSDB; AAP60578.
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31-JUL-1991
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AUGCCCAAGACCCAG---
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Db	Qy	Db	Qy	DЬ	Qy	Db	νQ
476 AAAGAUGGGCGUUUCCCCCAAUGAAAUACACAAGUAAACAU 516	478 AAAGAT-GGCATTTCCCCCAATGAAATACACAAGTAAACAT 517	416 UGCACGAGUUACCUGUUAAACUUUGGAACACCUACCAAAAAAAA	420 TGCAC-AGTTACCTG-TAAACATTGGAATACCGGCCAAAAAATAAGTTTGATCACATTTC 477	356 GAUGUAGGAAGACCCUCCUGAGGAGUGAAGAAGUGACAUGCCACCGCAGGAUCCUUUGCUC 415	361 GATGTA-GAAGACCCTTCTGAGGAGTGAAGGACGACGCACCGCAGGACCCTTTGCTC 419	300AAGGAAGUACAUUUGAAGAACGCAAGUAGAGGGAGUGCAGGAAACAAGAACUACAG 355	301 AGGAAAGGAAGTACATTTGAAGAACACAAGTAGAGGGAGTGCAGGAAACAAGAACTACAG 360

Search completed: June 15, Job time : 160.835 secs 2003, 16:08:47

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Scoring table:
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2: /cgn2_6/ptodata/2.
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/cgn2_6/ptodata/2/ina/5B_COMB.seq:*
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   GenCore version (c) 1993 - 2003
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PCT-US1-06452-1
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US-07-953-230A-1
US-07-953-230A-1
US-07-989-845-29
US-07-989-844-13
US-08-110-663-1
US-08-169-688-1
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US-08-169-686-1
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US-09-142-583A-10
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Sequence 1
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Patent No.
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Patent No.
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Patent No. 5405942
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Patent No. 5405942
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US-09-142-583A-3
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 Query Match
                                                                                                                                                                                                                  APPLICATION NUMBER: WO PCT/GB97/00658
FILING DATE: 11-MAR-1997
APPLICATION NUMBER: GB 9605124.8
FILING DATE: 11-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: SADOFF, B. J.
REGISTRATION NUMBER: 36663
REGISTRATION NUMBER: 117-263
TELEPHONE: 7038164000
TELEPHONE: 7038164100
                                                                                              FEATURE:
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Result

Database

Maximum

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## ALIGNMENTS

CORRESPONDENCE ADDRESS:
ADDRESSEE: NIXON &
STREET: 1100 NORTH TITLE OF INVENTION: METHOD OF TREATING NUMBER OF SEQUENCES: 11 APPLICANT: GOLDSPINK, GEOFFREY VANDERHYE P.C MUSCULAR DISORDERS

COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

OPERATING SYSTEM: PC-DOS/MS-DOS USA

APPLICATION NUMBER: US/09/142,583A
FILING DATE: 29-Oct-1998
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA: SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA: Version #1.25

INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 553 base pairs TYPE: nucleic acid STRANDEDNESS: both

MOLECULE TYPE: cDNA CDS 1..363 linear

18 19 20 21 23 24 25 27

ID NO:

ω

90.4%; Score 467.4; DB 4; Length 553;

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US-09-142-583A-5
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GENERAL INFORMATION:
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                                                                                                                              APPLICATION NUMBER: US/09/142,583A
FILING DATE: 29-Oct-1998
CLASSIFICATION: <UNKNOWN>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                             ZIP: 22201
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSE: NIXON & VANDERHYE P.C.
STREET: 1100 NORTH GLEBE ROAD
CITY: ARLINGTON
STATE: VA
                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: GOLDSPINK, GEOFFREY TITLE OF INVENTION: METHOD OF TREATING
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                                                              APPLICATION NUMBER: WO PCT/GB97/00658
FILING DATE: 11-MAR-1997
APPLICATION NUMBER: GB 9605124.8
FILING DATE: 11-MAR-1996
               NAME: SADOFF, B. J. REGISTRATION NUMBER: 36663
                                                                                                                                                                                                                                                                                                              COUNTRY: USA
REFERENCE/DOCKET NUMBER: 117-263
                                                                                                                                                                                                                                                                                                                                                                                                              OF SEQUENCES: 11
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US-09-142-583A-10
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                                                                                                                                                                         Sequence 10, Application US/09142583A Patent No. 6221842
GENERAL INFORMATION:
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TELEPHONE: 7038164000
TELEPAX: 7038164100
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 553 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
TRANDENESS: both
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                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: NIXON & VANDERHYE P.C
STREET: 1100 NORTH GLEBE ROAD
                                                                                                                          APPLICANT: GOLDSPINK, GEOFF
TITLE OF INVENTION: METHOD
NUMBER OF SEQUENCES: 11
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LOCATION: 341..397
SEQUENCE DESCRIPTION: S
COMPUTER READABLE FORM
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                                            CITY: ARLINGTON STATE: VA
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                               COUNTRY:
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                                                                                                                                            GEOFFREY
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Pred. No. 1.2e
0; Mismatches
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NAME/KEY: CDS:
COCATION: 26..493
SEQUENCE DESCRIPTION: SEQ ID NO:
US-09-142-583A-10
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SEQUENCE CHARACTERISTICS:
LENGTH: 777 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: WO PCT/GB97/00658
FILING DATE: 11 MAR-1997
APPLICATION NUMBER: GB 9605124.8
FILING DATE: 11-MAR-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: 7038164000
TELEFAX: 7038164100
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APPLICATION NUMBER: US/09/142,583A
FILING DATE: 29-Oct-1998
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
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                 AAAGAT-GGCATTTCCCCCAATGAAATACACAAAGTAAAACAT 517
                                                                                                                           GATGTAGGAAGACCCTCCTGAGGAGTGAAGAAGTGACATGCCACCGCAGGATCCTTTGCTC
                                                                                                                                                      GATGTA-GAAGACCCTTCTGAGGAGTGAAGAAGGACAGGCCACCGCAGGACCCTTTGCTC 419
                                                                                                                                                                                                          ATGCCCAAGACCCAGAAGTATCAGCCCCCATCTACCAACAAGAACACGAAGTCTCAGAGA
                                                                                                                                                                                                                                                                                                                                       TGCGCACCCTCAAGCCTGCCAAGTCAGCTCGCTCTGTCCGTGCCCAGCGGCCACACCCGAC
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AAAGATGGGCGTTTCCCCCAATGAAATACACAAGTAAACAT
                                                             TGCACGAGTTACCTGTTAAACTTTGGAACACCTACCAAAAAATAAGTTTGATAACATTTA
                                                                                          TGCAC-AGTTACCTG-TAAACATTGGAATACCGGCCCAAAAATAAGTTTGATCACATTTC
                                                                                                                                                                                                                                                                                                                        TGCGCACCCCTCAAGCCTGCCAAGTCAGCTCGCTCTGTCCGTGCCCCAGCGCCCACACCGAC
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                                                                                                                                                                                                                                                            ATGCCCAAGACCCAG-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: SADOFF, B. J.
REGISTRATION NUMBER: 36663
REFERENCE/DOCKET NUMBER: 117-263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                          AAGGAAGTACATTTGAAGAACGCAAGTAGAGGGAGTGCAGGAAACAAGAACTACAG
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87.1%;
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Pred. No. 3.
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RESULT 5 US-08-472-809B-8

Sequence 8, Application US/08472809B Patent No. 5925564 GENERAL INFORMATION:

APPLICANT: Schwartz, Robert J.
APPLICANT: DeMayo, Franco J.
APPLICANT: O'Mailey, Bert W.
TITLE OF INVENTION: Expression Vector Systems
TITLE OF INVENTION: Method of Use
NUMBER OF SEQUENCES: 8

CORRESPONDENCE ADDRESS

Lyon & Lyon

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TITLE OF INVENTION: PREPRO INSULIN-LIKE GROWTH FACTORS; I AND II
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO: 2
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Best Local
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 630,557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA: APPLICATION NUMBER: US/07/65,673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: BELL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH:
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                   AAAGAT-GGCATTTCCCCCAATGAAATACACAAGTAAACAT 517
                                                                               TGCAC-AGTTACCTG-TAAACATTGGAATACCGGCCAAAAAATAAGTTTGATCACATTTC
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                                                                                                                                             GATGTA-GAAGACCCTTCTGAGGAGTGAAGAAGGACAGGCCACCGCAGGACCCTTTGCTC 419
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AAAGAUGGGCGUUUCCCCCCAAUGAAAUACACAAGUAAACAU 516
                                                               UGCACGAGUUACCUGUUAAACUUUGGAACACCUACCAAAAAAUAAGUUUGAUAACAUUUA
                                                                                                                           GAUGUAGGAAGACCCUCCUGAGGAGUGAAGAGUGACAUGCCACCGCAGGAUCCUUUGCUC
                                                                                                                                                                                                                        AGGAAAGGAAGTACATTTGAAGAACACAAGTAGAGGGAGTGCAGGAAACAAGAACTACAG
                                                                                                                                                                                                                                                                                  ATGCCCAAGACCCCAGAAGTATCAGCCCCCCATCTACCAACAAGAACACGAAGTCTCAGAGA
                                                                                                                                                                                                                                                                                                                    UGCGCACCCCUCAAGCCUGCCAAGUCAGCUCGCUCUGUCCGUGCCCAGCGCCACACCGAC
                                                                                                                                                                                                                                                                                                                                             TGCGCACCCCTCAAGCCTGCCAAGTCAGCTCGCTGTCCGTGCCCAGCGCGCCACACCGAC
                                                                                                                                                                                       ----AAGGAAGUACAUUUGAAGAACGCAAGUAGAAGGGAGUGCAGGAAACAAGAACUACAG
                                                                                                                                                                                                                                                        AUGCCCAAGACCCAG----
                                                                                                                                                                                                                                                                                                                                                                              ACAGGUAUCGUGGAUGAGUGCUGCUUCCGGAGCUGUGAUCAUAGGAGGCUGGAGAUGUAU
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Pred. No. 3.4e-95
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Best Local
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REFERENCE/DOCKET NUMBER: 214/.

TELECOMMUNICATION INFORMATION:

TELEPHONE: (213) 489-1600

TELEFAX: (213) 955-0440

TELEX: 67-3510

INFORMATION TOT
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09,
FILING DATE: March 9,
APPLICATION NUMBER: 07
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 5707 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
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MOLECULE TYPE:
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ATTORNEY/AGENT INFORMATION:
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OPERATING SYSTEM: IBM F.C.
OPERATING SYSTEM: IBM F.C.
OPERATION DATA:
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                                                        1104
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                                                                                      361
                                                                                                                                301 AGGAAAGGAAGTACATTTGAAGAACACAAGTAGAAGGGAGTGCAGGAAACAAGAACTACAG
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California
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                                                                            GATGTA -GAAGACCCTTCTGAGGAGTGAAGAAGGACAGGCCACCGCAGGACCCTTTGCTC
                             TGCA 423
                                                          GATGTAGGAAGACCCTCCTGAGGAGTGAAGAGTGACATGCCACCGCAGGATCCCCCGGGC
                                                                                                                 ----AAGGAAGTACATTTGAAGAACGCAAGTAGAGGGAGTGCAGGAAACAAGAACTACAG
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07/789,919
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Pred. No. 1.9e-78;
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Best Local S
Matches 363
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/472,809B
FILING DATE: June 7, 1995
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: 08/209,846
APPLICATION NUMBER: 08/209,846
FILING DATE: March 9, 1994
APPLICATION UNMBER: 07/789,919
FILING DATE: NO. 5925564ember 6, 19
ATTORNEY/AGENT INFORMATION:
NAME: WEATHING BIORATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (213) 489-:
TELEFAX: (213) 955-04-
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 6345 bases
                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (213) 489-1600
TELEPHONE: (213) 955-0440
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 633 West Fit
STREET: Suite 4700
CITY: Los Angeles
STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITLE OF INVENTION:
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OPERATING SYSTEM:
                                                                                                                                                                                                                                                         / Match 55.4%;
Local Similarity 85.6%;
les 363; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 214/212
                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Warburg, Richard J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
                                             3822
                                                                                                                  3762
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                                                                                                                                                  61 AGGGGCTTTTATTTCAACAAGCCCACAGGGTATGGCTCCAGCAGTCGGAGGGCGCCTCAG
                                                                                                                                                                                                                                                                                                                                                                                                    nucleic acid
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TGCGCACCCCTCAAGCCTGCCAAGTCAGCTCGCTCTGTCCGTGCCCAGCGCCACACCCGAC
                                                                                ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTAT
                                             ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTAT
                                                                                                                  AGGGGCTTTTATTTCAACAAGCCCACAGGGTATGGCTCCAGCAGTCGGAGGGCCCCCCAG
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O'Malley, Bert W.
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RESULT 7
5405942-13
Patent No. 5405942
PAPPLICANT: BELL, G
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5405942-9
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TITLE OF INVENTION: PREPRO INSULIN-LIKE GROWTH FACTORS
                                                                                                                                    JAMES P
                                                                                                                                                                  Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO:1
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Best Local Similarity
Matches 257; Conserv
                                                                                                        AND II
   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/65,673
FILING DATE: 16-JUN-1987
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 630,557
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APPLICATION NUMBER: 630,557
FILING DATE: 19-JUL-1984
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CURRENT APPLICATION NUMBER: U$/07/65,673
APPLICATION NUMBER: 150-701-1987
                                                                                                                  TITLE OF INVENTION: PREPRO INSULIN-LIKE GROWTH FACTORS
                                                                                     NUMBER OF SEQUENCES: 16
                                                                                                                                                 APPLICANT: BELL,
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RESULT 10
5405942-11
;Patent No. 5405942
; APPLICANT: BELL, (
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LENGTH: 357
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Best Local Similarity
Matches 163; Conserv
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Best Local Similarity
Matches 206; Conserv
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APPLICATION NUMBER: 630,557
FILING DATE: 19-JUL-1984
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 16
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/65,673
FILING DATE: 16-JUN-1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: PREPRO INSULIN-LIKE GROWTH FACTORS
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                                                                                                 UGCGCACCCCUCAAGCCUGCCAAGUCAGCU
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     GRAEME I.; RALL,
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79.2%; Pred. No. 7.1e-69;
htive 50; Mismatches 4
     LESLIE
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     B.; MERRYWEATHER
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US-09-255-829-13
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SEQ ID NO:11:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local (
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MEDIDIM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/07/
FILING DATE: 16-JUN-1987
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 630,557
                                                                                                                                                  APPLICATION NUMBER: US/09/255,829
FILING DATE: 23-FEB-1999
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/GB97/02273
FILING DATE: 22-AUG-1997
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                             SOFTWARE: Patentin Recoursely Application Data
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                                                                                                   APPLICATION NUMBER: US 0: FILING DATE: 27-DEC-1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 16
CURRENT APPLICATION DATA:
                               REGISTRATION NUMBER: 32,893
REFERENCE/DOCKET NUMBER: 15
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Shone, Clifford Charles APPLICANT: Quinn, Conrad Padraig APPLICANT: Foster, Keith Alan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: PREPRO INSULIN-LIKE GROWTH FACTORS
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                TELEPHONE:
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202-371-2540
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27-DEC-1996
                                                                                    ROBERT W
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Pred. No. 4.8e-55;
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E, NW, SUITE 600
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US-09-255-829-13
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Best Local S
Matches 162
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Best Local
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                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/65,673
FILING DATE: 16-JUN-1987
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 630,557
                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: PREPRO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: BELL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
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 UGCGCACCCCUCAGGCCUGCCAAGUCAGCU
                            TGCGCACCCCTCAAGCCTGCCAAGTCAGCT 210
                                                        ACAGGUAUCGUGGAUGAGUGCUGCUGCGGAGCUGUGAUCUAAGGAGGCUGGAGAUGUAU
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Pred. No. 1.
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Pred. No. 1.5e-54;
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ZIP: 92101-2920

ZIP: 92101-2920

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

MEDIUM TYPE: IBM PC compatible

TOMPITER: IBM PC compatible

TOMPITER: PC-DOS/MS-DOS

TOMPITER: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (619)238-0999
TELEFAX: (619)238-0062
NFORMATION FOR SEQ ID NO: 1:
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APPLICANT:
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SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 240 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 07/983,523
FILING DATE: 03-MAR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/578,728
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 04-SEP-1990 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 51
FELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ITLE OF INVENTION: Production of Insulin-Like Growth
ITLE OF INVENTION: Factor-1 in Methylotrophic Yeast Cells
                                                                                                                                                                                                                                                                                                          Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: double
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: California
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                      TGCGCACCCCTCAAGCCTGCCAAGTCAGCTCGCT 214
                                                                                           ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTAT 180
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TGCGCACCCCTCAAGCCTGCCAAGTCAGCTTGAT 230
                                                                      ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTCGAGATGTAT 196
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1660 Union Street
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Howard, Bradley D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Davis, Geneva R.
Holtz, Gregory C.
                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     unknown
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09-SEPT-1994
                                                                                                                                                                                                                                                                                                        39.28;
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Pred. No. 2.7e-53;
0; Mismatches 7
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: PCT/US91/06452
FILING DATE: 19910409
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/578,728
FILING DATE: 04-SEP-1990
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie L.
NAME: Seidman, Stephanie L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (619)552-1311
TELEFAX: (619)552-0095
INFORMATION FOR SEQ ID NO: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Z1F: UNIVERSE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC -DOS/MS-DOS
COMPRATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 240 base pairs
TYPE: NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 51874
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619)552-1311
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 135 Street CITY: Chicago
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THE OF INVENTION:
THE OF SEQUENCES:
                                                                     137
197
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                                                                                                                                                                61 AGGGGCTTTTATTTCAACAAGCCCCACAGGGTATGGCTCCAGCAGTCGGAGGGCGCCTCAG
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                   TGCGCACCCCTCAAGCCTGCCAAGTCAGCTCGCT 214
                                                                                          ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTAT
                                                                                                                                          AGGGGCTTTTATTTCAACAAGCCCACAGGGTATGGCTCCAGCAGTCGACGGGCGCCTCAG
                                                                     ACAGGCATCGTGGATGATGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTCGAGATGTAT
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                                                                                                                                                                                                                                                                                           Conservative
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Factor-1 in Methylotrophic Yeast Cells
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Pred. No. 2.7e-53;
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RESULT 15 US-09-029-267-13

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TOPOLOGY: linear
TOPOLOGY: linear
HOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "Synthetic"
US-09-029-267-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/029,267
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
NAME: Chung, Ling-Fong
REGISTRATION NUMBER: 36,482
REFERENCE/DOCKET NUMBER: 1165.100
TELECOMMUNICATION INFORMATION:
TELEPHONE: (510) 601-2704
TELEPAX: (510) 655-3542
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARRCTERISTICS:
LENGTH: 390 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: 110-27
                                                                                                                                                                                                                                                                                                                                           Query Match 39.2%;
Best Local Similarity 96.7%;
Matches 207; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Zaror, Isabel
APPLICANT: Innis, Michael
TITLE OF INVENTION: Pichia Secretory Leader for Protein
TITLE OF INVENTION: Expression
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Emeryville
STATE: California
COUNTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET:
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                      181 TGCGCACCCCTCAAGCCTGCCAAGTCAGCTCGCT 214
                                                                                                             121 ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTAT 180
    340
                                                                                                                                                                     220 AGGGGCTTTTATTTCAACAAGCCCACAGGGTATGGCTCCAGCAGTCGACGGGCGCCTCAG
                                                                                                                                                                                                                                                       160 GGACCGGAGACGCTCTGCGGGGCTGAGCTCGTGGATGCTCTGCAGTTCGTGTGGAGAC 219
                                                                                    280 ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTCGAGATGTAT 339
                                                                                                                                                                                            61 AGGGGCTTTTATTTCAACAAGCCCACAGGGTATGGCTCCAGCAGTCGGAGGGGCGCCTCAG 120
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TGCGCACCCCTCAAGCCTGCCAAGTCAGCTTGAT 373
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                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                           Score 202.8; DB 3; Pred. No. 3.3e-53; 0; Mismatches 7;
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Search completed: June 15, 2003, 18:20:40 Job time : 43.9101 secs

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Minimum
Maximum
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Maximum Match 100%
Listing first 45 summaries
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length: 2000000000
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Match
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1: /cgn2_6/ptodata/2/p:
2: /cgn2_6/ptodata/2/p:
3: /cgn2_6/ptodata/2/p:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*
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/cgn2_6/ptodata/2/pubpna/US09_RUBCOMB.seq:*
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/cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US10_PUBCOMB.seq:*
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6/ptodata/2/pubpna/PCTUS_PUBCOMB.seq:*
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10 US-09-810-497-24
10 US-09-810-261-3
10 US-09-852-261-3
10 US-09-852-261-11
10 US-09-852-261-11
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10 US-09-852-261-11
10 US-09-852-261-11
10 US-09-852-261-13
10 US-10-161-088-3
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10 US-10-77-381-9
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Compugen Ltd.
                      Sequence 4, Appli
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Sequence 373, Appli
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Sequence 3, Appli
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Sequence 5, Appli
Sequence 13, Appl
Sequence 40,
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CURRENT FILING DATE: 2001-05-10
PRIOR APPLICATION NUMBER: GB 0011278.9
PRIOR FILING DATE: 2000-05-10
NUMBER: OF SEQ ID NOS: 14
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1
LENGTH: 517
TYPE: DNA
ORGANISM: Himmo sapiens
US-09-852-261-1
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RESULT 1 US-09-852-261-1

Sequence 1, Application US/09852261
Patent No. US20020083477A1
GENERAL INFORMATION:
APPLICANT: GOLDSPINK, GEOFFREY
APPLICANT: TERENGH, GIORGIO
TITLE OF INVENTION: REPAIR OF NERVE DAMAGE
FILE REFERENCE: 117-351

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ALIGNMENTS	US-10-280-826-38 US-09-961-398-38 US-09-961-398-38 US-09-954-531-98 US-10-136-841-3 US-10-136-841-3 US-10-166-543-997 US-10-166-543-997 US-10-166-543-997 US-10-166-543-11040 US-10-166-543-428 US-10-166-543-428 US-10-166-543-428 US-10-166-543-41-1 US-10-166-543-1918 US-10-166-543-1918 US-10-166-543-119-18 US-10-166-543-119-37	
	Sequence 38, Appl Sequence 2082, App Sequence 39, App Sequence 917, App Sequence 428, App Sequence 1, Appli Sequence 1137, App Sequence 218, App Sequence 1137, App Sequence 1137, App Sequence 218, App Sequence 237, Appl Sequence 37, Appl Sequence 293, App Sequence 293, App Sequence 293, App Sequence 277, Appl	

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Query Match Best Local :

Similarity

100.0%;

Score 517; DB 10; Pred. No. 1.2e-159;

Length 517;

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RESULT 2
US-09-852-261-5
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APPLICANT: GOLDSPIKK, GEOFFRBY
APPLICANT: TERENGHI, GIORGIO
TITLE OF INVENTION: REPAIR OF NERVE DAMAGE
FILE REFERENCE: 117-351
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SEQ ID NO 5
LENGTH: 523
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Best Local Similarity 96.2%;
Matches 501; Conservative
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CURRENT FILING DATE: 2001-05-10
PRIOR APPLICATION NUMBER: GB 0011278.9
PRIOR FILING DATE: 2000-05-10
NUMBER OF SEQ ID NOS: 14
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                CAGGATGTA-GAAGACCCTTCTGAGGAGTGAAGAAGGACAGGCCACCGCAGGACCCTTTG
                                                                                                                                             ATGCCCAAGACCCAGAAGTATCAGCCCCCATCTACCAACAAGAACACGAAGTCTCA---G
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                                                            AGAAGGAAAGGAAGTACATTTGAAGAACACAAGTAGAGGGAGTGCAGGAAACAAGAACTA
                                                                                                                         ATGCCCAAGACTCAGAAGTATCAGCCTCCATCTACCAACAAGAAAATGAAGTCTCAGAGG
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Pred. No. 2.5e-143;
0; Mismatches 16;
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CURRENT APPLICATION NUMBER: US/09/852,261
CURRENT FILING DATE: 2001-05-10
PRIOR APPLICATION NUMBER: GB 0011278.9
PRIOR FILING DATE: 2000-05-10
NUMBER OF SEQ ID NOS: 14
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 13
LENGTH: 471
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Patent No. US20020083477A1
GENERAL INFORMATION:
APPLICANT: GOLDSBINK, GEOFFREY
APPLICANT: TERBURHI, GIORGIO
TITLE OF INVENTION: REPAIR OF NERVE D
FILE REFERENCE: 117-351
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US-09-852-261-13
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AGATGGCATTTCCCCCCAATGAAATACACAAGTAAACAT
                                                          TGCACAGTTACCTGTAAACATTGGAATACCGGCCAAAAAATAAGTTTGATCACATTTCAA
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RESULT 4 US-10-136-639-4

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RESULT 5
US-09-919-497-24
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                                                                                                                             Sequence 24, Application US/09919497
Patent No. US20020106662A1
GENERAL INFORMATION:
APPLICANT: Mutter, George L.
TITLE OF INVENTION: PROGNOSTIC CLASSIFICATION
FILE REFERENCE: B0801/7225
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LENGTH: 7260
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CURRENT FILING DATE: 2002-09-06
PRIOR APPLICATION NUMBER: US 60/329,650
PRIOR FILING DATE: 2001-10-16
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin version 3.0
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                    CURRENT APPLICATION NUMBER: US/09/919,497
CURRENT FILING DATE: 2001-07-31
PRIOR APPLICATION NUMBER: US 60/221,735
PRIOR FILING DATE: 2000-07-31
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OF SEQ ID NOS:
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87.3%;
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Pred. No. 2.8e-102;
0; Mismatches 13;
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                                                                                                                                                                ENDOMETRIAL CANCER
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; TYPE: DNA
; ORGANISM: HOMO s
US-09-919-497-24
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                                                                                                                                                                                  APPLICANT: Horne, Darci T.
APPLICANT: Vockley, Joseph G.
APPLICANT: Vockley, Joseph G.
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in FILL REFERENCE: 44921-5028-WO
FULL REFERENCE: 44921-5028-WO

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 3739, Application US/09880107 Patent No. US20020142981A1 GENERAL INFORMATION:
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SEQ ID NO 24
LENGTH: 7260
                                                                                                      SOFTWARE: PatentIn Ver.
SEQ ID NO 3739
LENGTH: 7260
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                                              TYPE: DNA
ORGANISM: Homo sapiens
  OTHER INFORMATION: Genbank Accession
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Pred. No. 2.8e
0; Mismatches
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RESULT 7
US-09-852-261-3
Sequence 3, Application US/09852261
Patent No. US20020083477A1
GENERAL INFORMATION:
APPLICANT: GOLDSPINK, GEOFFREY
APPLICANT: TREENING GOORGIO
TITLE OF INVENTION: REPAIR OF NERVE DAMAGE
FILE REFERENCE: 117-351
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                                                                                       Best Local
                                                                                                  Query Match
                                                                                                                                                                             SEQ ID NO 3
                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/852,261
CURRENT FILING DATE: 2001-05-10
                                                                                                                                                                                                     CURRENT FILING DATE: 2001-05-10
PRIOR APPLICATION NUMBER: GB 0011278.9
PRIOR FILING DATE: 2000-05-10
NUMBER OF SEQ ID NOS: 14
                                                                                                                                                                                         SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                 LENGTH: 539
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                                                                           similarity 81.3
29; Conservative
                        TGCAC-AGTTACCTG-TAAACATTGGAATACCGGCCAAAAAATAAGTTTGATCACATTTC 477
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 AGGGGCTTTTATTTCAACAAGCCCACAGGGTATGGCTCCAGCAGTCGGAGGGCGCCTCAG
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87.3%;
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Pred. No. 2.8e-102;
                                                                         Score 325.2; DB 10;
Pred. No. 1.4e-96;
D; Mismatches 88;
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Best Local Similarity
Matches 419; Conserv
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TGCGCACCCCTCAAGCCTGCCAAGTCAGCTCGCTCTGTCCGTGCCCAGCGCCCACACCGAC
                           ACAGGCATTGTGGATGAGTGTTGCTTCCGGAGCTGTGATCTGAGGAGACTGGAGATGTAC 318
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241 ATGCCCAAGACTCAGAAGTCCCAGCCCCTATCGACACAAGAAAAAGGAAAGCTGCAAAGG
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                                ACATITCAAAGAT-GGCATTTCCCCCCAATGAAATACACAAGTAAACAT 517
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TCATTTCAGAGATGGGCATTTCCCCTCAATGAAATACACAAGTAAACAT
                                                                                                                        CTGCTTGAGCAACCTGCAAAACATCGGAACACCTGCCAAATATCAATAATGAGTTCAATA 480
                                                                                                                                                                   CTCTGCACAGTTACCTGTAAACATTGGAATACCGGCCA-----AAAAATAAGTTTGATC
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Sequence 1, Application US/10161088
PUDLICATION NO. US20030077761A1
GENERAL INFORMATION:
APPLICANT: PAITOW, Vendela
APPLICANT: ROSENGIEN, LINDA
TITLE OF INVENTION: NEW METHODS
FILE REFERENCE: 13425-111001
CURRENT FAPLICATION NUMBER: US/10/161,088
CURRENT FILING DATE: 2002-05-31
PRIOR APPLICATION NUMBER: SE 0101934-8
PRIOR FILING DATE: 2001-06-01
NUMBER OF SEQ ID NOS: 3
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 651
TYPE: DWA
ORGANISM: Homo Sapiens AGGGGCTTTTATTTCAACAAGCCCACAGGGTATGGCTCCAGCAGTCGGAGGGGCCCTCAG 120 AGGGGCTTTTACTTCAACAAGCCCACAGGCTATGGCTCCAGCATTCGGAGGGCACCTCAG GGACCAGAGACCCTTTGCGGGGGCTGAGCTGGTGGATGCTCTTCAGTTCGTGTGTGGACCG 198 GGACCGGAGACGCTCTGCGGGGCTGAGCTGGTGGATGCTCTTCAGTTCGTGTGTGGAGAC 60 Score 318.2; DB Pred. No. 3e-94; 0; Mismatches DB 9; 83;

Indels Length

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Gaps

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US-09-852-261-11
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Best Local S
Matches 259
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PRIOR FILING DATE: 2000-05-10
NUMBER OF SEQ ID NOS: 14
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 9
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Sequence 11,
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CURRENT APPLICATION NUMBER: US/09/852,261
CURRENT FILING DATE: 2001-05-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: GOLDSPINK, GEOFFREY
APPLICANT: TERENGHI, GIORGIO
TITLE OF INVENTION: REPAIR OF NERVE DAMAGE
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                                                                            ATGCCCAAGACCCAGAAGGA
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Application US/0985226:
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99.6%;
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Pred. No. 9.5e-75;
0; Mismatches 1;
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Sequence 1, Application US/10238114
Publication No. US20030100073A1

GENERAL INFORMATION:
APPLICANT: MARTION:
APPLICANT: ANDREONI, Christine Michele
TITLE OF INVENTION: IGF-1 AS FELINE VACCINE AD:
FILE REPERENCE: 454313-3165.1

CURRENT APPLICATION NUMBER: US/10/238,114

CURRENT APPLICATION NUMBER: US/10/238,114

CURRENT APPLICATION NUMBER: US/10/238,114

PRIOR APPLICATION NUMBER: US/10/318,666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TYPE: DNA; ORGANISM: Rattus US-09-852-261-11
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SEQ ID NO 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/852,261
CURRENT FILING DATE: 2001-05-10
PRIOR APPLICATION NUMBER: GB 0011278.9
PRIOR FILING DATE: 2000-05-10
NUMBER OF SEQ ID NOS: 14
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APPLICANT: TERENGHI, GIORGIO
TITLE OF INVENTION: REPAIR OF NERVE DAMAGE
FILE REFERENCE: 117-351
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Pred. No. 3.6
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US-09-930-377B-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Wheeler, Matthew B.
APPLICANT: Donovan, Sharon M.
APPLICANT: Bleck, Gregory T.
APPLICANT: Monaco-Geigel, Marcia
TITLE OF INVENTION: Animals Expressing Exogenous IGF-I in their Milk
FILE REFERENCE: 66-00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 240; Conserv
                                                                                                                                                                              Best Local Similarity Matches 210; Conserv
                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 1
                                                                                                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/930,377B CURRENT FILING DATE: 2001-08-15 PRIOR APPLICATION NUMBER: 60/225,474 PRIOR FILING DATE: 2000-08-15 NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 2001-09-12
NUMBER OF SEQ ID NOS: 20
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                             TYPE: DNA ORGANISM: Artificial Sequence
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                                    AGGGGATTTTATTTCAACAAGCCCCACAGGGTATGGATCCAGCAGTCGGAGGGCGCCCCAG
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ACAGGCATCGTGGATGAGTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTAT
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Pred. No. 1.7e-56;
D; Mismatches 9;
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Pred. No. 1.1e-64;
0; Mismatches 20
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US-10-179-046-13
; Sequence 13, Application US/10179046
; Publication No. US20030013154A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/930,377B CURRENT FILING DATE: 2001-08-15 PRIOR APPLICATION NUMBER: 60/225,474 PRIOR FILING DATE: 2000-08-15 NUMBER OF SEO ID NOS: 2 SOFTWARE: Patentin Ver. 2.0 SEO ID NO 2 LENGTH: 210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 206; Conserv
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APPLICANT: Donovan, Sharon M.
APPLICANT: Bleck, Gregory T.
APPLICANT: Monaco-Seigel, Marcia
TITLE OF INVENTION: Animals Expressing Exogenous IGF-I in their
FILE REFERENCE: 66-00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA ORGANISM: Artificial Sequence
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                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Chiron
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COMPUTER READABLE FORM:
                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                  TITLE OF INVENTION: Pichia Secretory Leader for Protein
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                                  ADDRESSEE: Chiron Corporation
STREET: 4560 Horton Street
CITY: Emeryville
STATE: California
COUNTRY: United States
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Innis, Michael
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pred. No. 8e-57;
0; Mismatches 4;
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CURRENT APPLICATION NUMBER: US/10/161,088
CURRENT FILING DATE: 2002-05-31
PRIOR APPLICATION NUMBER: SD 0101934-8
PRIOR FILING DATE: 2001-06-01
NUMBER OF SEQ ID NOS: 3
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3
LENGTH: 286
TYPE: DNA
ORGANISM: Homo sapiens
US-10-161-088-3
                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 15
US-10-161-088-3
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RILING DATE: <UNKNOWN')

ATTORNEY/AGENT INFORMATION:

NAME: Chung, Ling-Fong

REGISTRATION NUMBER: 36,482

REFERENCE/DOCKET NUMBER: 1165.100

TELECOMMUNICATION UNFORMATION:

TELEPHONE: (510) 601-2704

TELEFAX: (510) 655-3542

INFORMATION FOR SEQ ID NO: 13:

SEQUENCE CHARACTERISTICS:
LENGTH: 390 base pairs

TYPE: nucleic acid

STRANDEDNESS: single

TOPCLOST: linear

MOLECULE TYPE: other nucleic acid

DESCRIPTION: /desc = "Synthetic"

SEQUENCE DESCRIPTION: SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                            Sequence 3, Application US/10161088 Publication No. US20030077761A1 GENERAL INFORMATION:
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Best Local Similarity
                      Query Match
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  Best Local Similarity
                                                                                                                                                                                                                                                                                      APPLICANT: Parrow, Vendela
APPLICANT: Rosengren, Linda
TITLE OF INVENTION: NEW METHODS
FILE REFERENCE: 13425-111001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TGCGCACCCCTCAAGCCTGCCAAGTCAGCTTGAT 373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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COMPUTER: IBM PC compati
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96.7%;
39.1%;
88.0%;
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Score 202; DB 9;
Pred. No. 3.1e-56;
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                   Length 286;
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258 GCATGCAAGC 267
                                     256 AAGTATCAGC 265
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Search completed: June 15, 2003, 20:22:23 Job time: 95.6428 secs

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Maximum DB seq length: 2000000000
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em_gss_pln:*
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DEFINITION
Clone IMAGE:1001007 3' similar to gh:N04482 Mouse mRNA for preproissulin-like growth factor IB (MOUSE); mRNA sequence.

ACCESSION
AI503976.1 GI:4401827
EST
SOURCE
ORGANISM
Mus musculus
ENdaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
Marrayota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
Marrayota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
Marrayota; Marrayota; Martin, J., Beck, C., Wylie, T.,
Underwood, K., Steptoe, M., Theising, B., Allen, M., Bowers, Y., Person, B., Swaller, T., Gibbons, M., Pape, D., Harvey, N., Schurk, R., Ritter
, E., Kohn, S., Shin, T., Jackson, Y., Cardenas, M., McCann, R.,
Waterston, R. and Wilson, R.
The Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108, USA
Tali 314 286 1810
Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LINL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.

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VERSION
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UI-CF-EC1-abj-k-24-0-UI 3', mRNA
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                                                                                                                                                                                                     CTCTGCACAGTTACCTGTAAACATTGGAATACCGGCCA------AAAAATAAGTTTGATC
                                                                                                                                                                                                                                     CAGAATGTAGGAGGAGCCTCCCACGGAGCAGAAAATGCCACATCACCGCAGGATCCTTTG
                                                                                                                                                                                                                                                  CAGGATGTA-GAAGACCCTTCTGAGGAGTGAAGGAAGGACAGGCCACCGCAGGACCCTTTG
                                                                                                                                                                                                                                                                                TECECACCCCTCAAGCCTECCAAGTCAGCTCGCTCTCTCTCCGTGCCCAGCGCCACACCGAC
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                                                                                                                                        ACATTACAAAGATGGGCATTTCCCCCCAATGAAATATACAAGTAAAACAT
                                                                                                                                                               ACATTTCAAAGAT-GGCATTTCCCCCCAATGAAATACACAAGTAAACAT 517
                                                                                                                                                                                       CTGCTTGAGCAACCTGCAAAACATCGAAACACCTACCAAATAACAATAATAAGTCCAATA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /tissue_type="diaphragm"
/dev_stage="adult"
/dev_stage="adult"
/lab_host="SOLR (kanamycin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone="IMAGE:1001007"
/clone_lib="Stratagene mouse diaphragm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Mus musculus"
/db_xref="taxon:10090"
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Pred. No. 1.4e-87;
0; Mismatches 84;
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I-CF-EC1 Homo sapiens
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                                                                                                                                                                                                  Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               McCray Lab
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            discovery
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2024 University of Tel: 319 356 4866 Fax: 319 356 7171
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Tissue Procurement: Dr. M. J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Genome Res. 6 (9), 97044477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 673)
Bonaldo,M.F., Lennon,G. and Soares,M.B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Genetics (www.resgen.com).
Seg primer: M13 FORWARD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tissue Procurement: Dr. M. J. Welsh, University of Iowa CDNA Library preparation: Dr. M. Bento Soares, University of Iowa CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa DNA Sequencing by: Dr. M. Bento Soares, University of Iowa Clone Distribution: Researchers may obtain clones from Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: McCray,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bonaldo, M.F., Lennon, G. and Soares.
Normalization and subtraction: two
ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTAT 180
                                                                                                                                                                                                                                                                                                   GACCGGAGACGCTCTGCGGGGCTGAGCTGGTGGATGCTCTTCAGTTCGTGTGTGGAGAC
                                                                                                                                                                                                                                                                                                                                GACCGGAGACGCTCTGCGGGGCTGAGCTGGTGGATGCTCTTCAGTTCGTGTGTGGAGAC
                                                                                                                                                                                                      AGGGG-TTTTATTTCAGCAAGCCCACAGGGTATGGCTCCAGCAGTCGGAGGGCCCCTCAG
                                                                                                                                                                                                                                   AGGGGCTTTTATTTCAACAAGCCCACAGGGTATGGCTCCAGCAGTCGGAGGGCGCCTCAG
                                                                                                    ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTAT
                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /dev_stage="Adult and Fetal"
//dev_stage="Adult and Fetal"
//lab host="DH108 (Life Technologies) (Tl phage resistant)"
//note="Organ: Lung: Vector: pT7T3-Pac (Pharmacia) with a
modified polylinker; Site_1: EcoR I; Site_2: Not I;
UI-CF-EC1 is a normalized cDNA library containing the
following tissue(s): Normal lung from adult and from fetal
day 64, day 87, week 19 and week 42. The library was
constructed according to Bonaldo, Lennon and Soares,
Genome Research, 6:791.806, 1996. First strand cDNA
synthesis was primed with an oligo-dT primer containing a
Not I site. Double stranded cDNA was ligated to an EcoR I
adaptor, digested with Not I, and cloned directionally
into pT7T3-Pac vector. The oligonuclectide used to prime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the synthesis of first-strand cDNA contains a library tag sequence that is located between the Not I site and the (dT)18 tail. The sequence tag for this library is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TAG_LIB=UI-CF-EC1
TAG_TISSUE=Normal Lung
and 380-383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/db_xref="taxon:9666"
/clone="UI-CF-EC1-abj-k-24-0-UI"
/clone_1lb="UI-CF-EC1"
/tissue_type="Tung"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TAG_SEQ=AAGTGCTTAC"
164 c 169 g
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Seq primer:
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Mus.

1 (bases 1 to 623)

1 (bases 1 to 623)

Marra, M., Hillier, L., Kucaba, T., Martin, J., Beck, C., Wylie, T.,

Underwood, K., Steptoe, M., Theising, B., Allen, M., Bowers, Y., Person, B., Swaller, T., Gibbons, M., Pape, D., Harvey, N., Schurk, R., Ritter, E., Kohn, S., Shin, T., Jackson, Y., Cardenas, M., McCann, R.,

Waterston, R. and Wilson, R.

The WashU-NCI Mouse EST Project 1999

Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AW146128. 623 bp mRNA linear EST 10-c um37e10.x1 Sugano mouse embryo mewa Mus musculus cDNA clone IMAGE:2247498 3' similar to gb:X04482 Mouse mRNA for preproinsulin-like growth factor IB (MOUSE);, mRNA sequence. AW146128 AW146128.1 GI:6167864 EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Marra M/WashU-NCI Mouse EST Project Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. Louis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL ;
IMAGE Consortium (info@image.llnl.gov) for further ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  house mouse
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314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                       quality sequence stop:
                                                                                                                                                                                                            /strain="C57BL"
/db_xref="taxon:10090"
/clone="IMAGE:2247498"
/clone_Lib="Sugano mouse embr
/dev_stage="embryo, 14 dpc"
/lab_host="DH10B"
                                   /note="Vector: pmE18S-FL3; Site_1: DraIII (CACTGTGTG); Site_2: DraIII (CACCATGTG); Ist strand cDNA was primed with an oligo(dT) primer (ATGTGCCCTTTTTTTTTTTTTTTT)

double-stranded cDNA was ligated to a DraIII adaptor [TGTTGCCCTACTGG], digested and cloned into distinct DraIII sites of the pmE18S-FL3 vector (5' site CACTGTGTG, 3' site CACCATGTGG). XhoI should be used to isolate the cDNA insert. Size selection was performed to exclude fragments
(University of Tokyo
                                                                                                                                                                                                                                                                                                                                                       /organism="Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                     Library constructed by Dr.
                                                                                                                                                                                                                                                                                                                                                                                                                         499.
Institute of Medical Science)
                                                                                                                                                                                                                                                                embryo mewa"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        further information.
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                     Sumio Sugano
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
                                     Contact: Robert Strausberg, Ph.D.
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL
IMAGE Consortium (info@image.llnl.gov) for further
Insert Length: 918 Std Error: 0.00
Seg primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                         AI248089 575 bp mRNA linear EST 01-DEC qh69f05.x1 Soares_fetal_liver_spleen_INFLS_S1 Homo sapiens cDEC clone IMAGE:1849953 3' similar to gb:X57025_rnal INSULIN-LIKE GROWTH FACTOR IA PRECURSOR (HUMAN);, mRNA sequence.
                                                                                                                                                                         Tumor Gene Index
Unpublished (1997)
                     High quality sequence stop: 380
                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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CTTCTGCTCTAAAAGCTGCG and 3' end
CGACCTGCAGCTCGAGCACA."
a 138 c 170 g 191 t
    Location/Qualifiers
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                                                                                                                                                                                                                                                    TCCTGAGGAGTGAAGAGTGACATGCCACCGCAGGATCCTTTGCTCTGCACGAGTTACCTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/db_xref="taxon:9666"
/clone="InAGE:1849953"
/clone=11b="Scares_fetal_liver_spleen_1NFLS_S1"
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/lab_host="DH10B (ampicillin resistant)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /sex="male"
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Mammalia; ]
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248 AGACCCAGAAGTATCAGCCCCCATCTACCAACAAGAACACGAAGTCTCA----GAGAAGGA
                                                                                                                                                                                              129 AGCAACCTGCAAAACATCGGAACACCTGCCAAATATCAATAATGAGTTCAATACCATTTC
                                                                                                                                                                                                                                     424 CAGTTACCTGTAAACATTGGAATACCGGCCA-----AAAAATAAGTTTGATCACATTTC
                                                                                                                                                                                                                                                                                 189 TAGGAGGAGCCTCCCGAGGAACAGAAAATGCCACGTCACCGCAAGATCCTTTGCTGCTTG
                                                                                                                                                                                                                                                                                                                        365 TA-GAAGACCCTTCTGAGGAGTGAAGAAGGACAGGCCACCGCAGGACCCTTTGCTCTGCA
                                                                                                                                                                                                                                                                                                                                                                   249 AAGGAAGTACACTTGAAGAACACAAGTAGAGGAAGTGCAGGAAACAAGACTTACAGAATG
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On Oct 6, 1998 this se
Other_ESTs: TC50779
Contact: Lee, NH
The Institute for Geno
9712, Medical Center D
Tel: (301)-838-3529
Fax: (301)-838-0208
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Rat Genome Project: Generation of a Rat EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CCCTCAAGCCTGCCAAGTCAGCTCGCTCTGTCCGTGCCCAGCGCCCACACCGACATGCCCA
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                                                                                                           AGAGATGGGCATTTCCCTCAATGAAATACACAAGTAAACAT
                                                                                                                                                  AAAGAT-GGCATTTCCCCCAATGAAATACACAAGTAAACAT 517
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         il: nhlee@tigr.org
primer: M13-21.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Rattus sp."
/db_xref="taxon:10118"
/clone="RKIBP33"
/clone=Ine"Normalized rat kidney, Bento Soares"
/note="Organ: kidney; Vector: pT7T3Pac; Site_1: EcoRI;
Site_2: NotI"
140 c 133 g 164 t
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REFERENCE
AUTHORS
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1 (bases 1 to 558)
Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrich,N., Dubuque,T. Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M., Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B., Theising,B., Wylie,T., Lennon,G., Soares,B., Wilson,R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            uj04b07.x1 Sugano mouse liver mlia Mus musculus cDNA clone IMAGE:1890901 3' similar to gb:X04482 Mouse mRNA for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: mouseestewatson.wustl.edu
This clone is available royalty-free through LLNL; contact the
INAGE Consortium (info@image.llnl.gov) for further information.
mGI:975225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WashU-HHMI Mouse EST Project
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       preproinsulin-like growth factor IB (MOUSE);, mRNA sequence
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (1996)
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                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                AGGGGCTTTTATTTCAACAAGCCCACAGGGTATGGCTCCAGCAGTCGGAGGGCGCCCCAG 120
                                         ACAGGCATTGTGGATGAGTGTTGCTTCCGGAGCTGTGATCTGAGGAGACTGGAGATGTAC
                                                                              ACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTAT 180
                                                                                                                                                                                                       GGACCAGAGACCCTTTGCGGGGCTGAGCTGGTGGATGCTCTTCAGTTCGTGTGGGACCG 447
                                                                                                                                                                                                                                              GGACCGGAGACGCTCTGCGGGGCTGAGCTGGTGGATGCTCTTCAGTTCGTGTGGAGAC
                                                                                                                                                                                                                                                                                                                                                                                             106
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                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                        /note="Organ: liver; Vector: pME18S-FL3; Site_1: DraIII (CACTGTGTG); Site_2: DraIII (CACCATGTG); lst strand cDNA was primed with an oligo(dT) primer
[ATGTGGCCTTTTTTTTTTTTTTTTT]; double-stranded cDNA was ligated to a DraIII adaptor [TGTTGGCCTACTGG], digested and cloned into distinct DraIII sites of the pME18S-FL3 vector (5' site CACTGTGTG, 3' site CACCATGTG). XhoI should be used to isolate the cDNA insert. Size selection was performed to exclude fragments <1.5kb. Library constructed by Dr. Sumio Sugano (University of Tokyo Institute of Medical Science). Custom primers for sequencing: 5' end primer CTGCTGCTCTAAAAGCTGCG and 3' end primer CGACCTGCACCTCACACA."
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/clone_lib="Sugano mouse liver mlia"
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/strain="C57BL"
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/lab_host="DH10B"
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80.8%;
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326 TGTGCCCCACTGAAGCCTACAAAAGCAGCCCGCTCTATCCGTGCCCAGCGCCACACTGAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          358 CAGGATGTA-GAAGACCCTTCTGAGGAGTGAAGAAGGACAGGCCACCGCAGGACCCTTTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cDNA Library Preparation: M. Bento Soares, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 603 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: cgapbs-rémail.nih.gov
Tissue Procurement: Michael J. Brownstein, M.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ni98cl0.sl NCI_CGAP_Pr21 Homo sapiens cDNA clone IMAGE:984882 3' similar to gb:X57025_rnal INSULIN-LIKE GROWIH FACTOR IA PRECURSOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seq primer: -40ml3 fwd. ET from Amersham High quality sequence stop: 412.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota: Metazoa: Chordata: Craniata; Vertebrata: Mammalia; Eutheria; Primates; Catarrhini; Hominidae: 1 (bases 1 to 498)
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National Cancer Institute, Cancer Genome Anat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (HUMAN);, mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                      /tissue_type="normal prostate"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                    /sex="male"
                                                                                                                                                                                                                                                                                                                                                                           /clone="IMAGE:984882"
/clone__lib="NCI_CGAP_Pr21"
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TITLE
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                                                                                                                                                                                                                                                                                                                                                                ue94h02.yl Sugano mouse embryo mewa Mus musculus cDNA clone IMAGE:1498803 5' similar to gb:X04482 Mouse mRNA for preproinsulin-like growth factor IB (MOUSE);, mRNA sequence. AII19218
AII19218.1 GI:3519542
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                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Verrebrata; Euteleostomi; Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 816)

Marra, M., Hillier, L., Allen, M., Bowles, M., Dietrich, N., Dubuque, T., Geisel, S., Kucaba, T., Lacy, M., Le, M., Martin, J., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Underwood, K., Moore, B., Theising, B., Wylle, T., Lennon, G., Soares, B., Wilson, R. and Waterston, R.
                                                  Unpublished (1996)
Contact: Marra MyMouse EST Project
Washbu-HHMI Mouse EST Project
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Lc
Tel: 314 286 1800
Fax: 314 286 1810
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l: mouseest@watson.wustl.edu clone is available royalty-free through LLNL; contact the Consortium (info@image.llnl.gov) for further information.
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86.2%;
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ACCESSION
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384; Conser
mRNA sequence.
BF383724
BF383724.1 GI:
EST.
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602044632F1 NCI_CGAP_Li9
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h quality sequence stop: 47
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="Vector: pME18S-FL3; Site_1: DraIII (CACTGTGTG);
Site_2: DraIII (CACCATGTG); 1st strand collaw, was primed
with an oligo(dT) primer (ATGTGGCTTTTTTTTTTTT);
double-stranded cDNA was ligated to a DraIII adaptor
[TGTTGGCCTACTGG], digested and cloned into distinct DraIII
sites of the pME18S-FL3 vector (5' site CACTGTGTG, 3' site
CACCATGTG). XhoI should be used to isolate the cDNA
insert. Size selection was performed to exclude fragments
<1.5kb. Library constructed by Dr. Sumio Sugano
(University of Tokyo Institute of Medical Science).
Custom primers for sequencing: 5' end primer
CGACCTGCAGCTGCAGCACA."

CGACCTGCAGCTGCAGCACA."

8 others
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/clone="IMAGE:1498803"
/clone_lib="Sugano mouse embryo mewa"
/dev_stage="embryo, 14 dpc"
/lab_host="DH10B"
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/strain="C57BL"
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                                                                      594
Mus
                                                                      musculus
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Anatomy Project (CGAP),

Euteleostomi,

contact th information

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AUTHORS
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Matches 394
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                                    486
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llni.gov
plate: LLAM9527 row: p column: 08
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1 (bases 1 to 594)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: cgapbs-r@mail.nih.gov
Tissue Frocurement: Jeffrey E. Green,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg, Ph.D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1999)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CISSUE Procurement: Jeffrey E. Green, M.D. CDNA Library Preparation: Life Technologies, Inc. CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                      ACATTTGAAGAACACAAGTAGAGGGAGGGAGTGCAGGAAACAAGAACTACAGGATGTA-GAAGA
                                                                                                                                                                                                                                                                                                                                                                                                                                       GAGTGTTGCTTCCGGAGCTGTGATCTGAGGAGACTGGAGATGTACTGTGCCCCCACTGAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTATTGCGCACCCCTCAAG
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                                    CATTICCC 493
                                                                                                                                                                 CCCTTCTGAGGAGTGAAGGACAGGCCACCGCAGGGACCCTTTGCTCTGCACAGTTACC
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GCATTGCC
                                                                       GCAAAACATCGAAACACCTACCAAATAACAATAATAAGTCCAATAACATTACAAAGATGG
                                                                                            TGTAAACATTGGAATACCGGCCA-----AAAAATAAGTTTGATCACATTTCAAAGATGG
                                                                                                                                              GCCTCCCACGGAGCAGAAAATGCCACATCACCGCAGGATCCTTTGCTGCTTGAGCAACCT
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                                                                                                                                                                                                                                                                                          musculus
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/db_xref "taxon:10090"
/dlone="LMAGE::1194295"
/clone=lib="NGT_CGAP_Li9"
/lab_host="DH10B (Tl phage-resistant)"
/note="Organ: liver; Vector: pcMV-SPORT6; Site_1: Not Site_2: Sall; cloned unidirectionally. Primer: Oligo Average insert size 1.9 kb. Constructed by Life Technologies. Note: this is a NCI_CGAP Library."
75 a 162 c 142 g 115 t
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Pred. No. 3e
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3e-79;
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nes 413; Conserv
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Old Service AA913900
AA913900.s2 Soares_NFL_T_GBC_1 Homo sapiens CDNA clone IMAGE:1525496 3' similar to gb:x57025_rnal INSULIN-LIKE GROWTH FACTOR IA PRECURSOR (HUMAN); mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mammalia; Eutheria;
l (bases 1 to 527)
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                                                                                                                                                                                                                                                                                                          TCTTCAGTTCGTGTGGAGACAGGGGCTTTATTTACAACAAGCCCACAGGGTATGGCTC
                                                                                                                                                                                                                                     CAGCAGTCGGAGGGCGCCTAAGACAGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGA
                                                                                                                                                                                                                                                        CAGCAGTCGGAGGCGCCTCAGACAGGCATCGTGGATGAGTGCTGCTTCCCGGAGCTGTGA
                                                                                                                                                                                                                                                                                                                                              TCTTCAGTTCGTGTGGGAGACAGGGGCTTTTATTTCAACAAGCCCACAGGGTATGGCTC
                                                   CAAGAACACGAAGTCTCAGAGAAGGAAAGGAAGTACATTTGAAGAACACAAGTAGAGGGA
                                                                                        CCGTGCCCAGCGCCACACCGACATGCCCAAGACCCAG
                                                                                                                                                                                  CCGTGCCCAGCGCCACACGACATGCCCAAGACCCAGAAGTATCAGCCCCCATCTACCAA
                                                                                                                                                               125
                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="Organ: pooled; Vector: pT773D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Egual amounts of plasmid DNA from three normalized libraries (fetal lung NbHL19W, testis NHT; and B-cell NCI_CGAP_GCB1) were mixed, and scircles were made in vitro. Following HAP purification reaction. The driver tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 lbraries. The pools consisted of I.M.A.G.E. clones 297480-302087, 682632-687239, 726408-728711, and 729096-731399. Subtraction by Bento Soarzes and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:1525496"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="Soares_NFL_T_GBC_S1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ocation/Qualifiers.
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                                                                                                                                                                                                                                                                                                                                                                                                  58.0%;
85.5%;
                                                                                                                                                                                                                                                                                                                                                                             Score 299.8; DB y;
Pred. No. 3.9e-78;
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Length Indels

527;

53,

Gaps

277

278

218 408 158 468 98

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RESULT 11
AI876493/c
LOCUS
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ORGANISM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1999)
Contact: Marra M/WashU-NCI Mouse EST Project 1999
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 642)
Marra, M., Hillier, L., Kucaba, T., Martin, J., Beck, C., Wylie, T., Underwood, K., Steptoe, M., Theising, B., Allen, M., Bowers, Y., Person, B., Swaller, T., Gibbons, M., Pape, D., Harvey, N., Schurk, R., Ritter, E., Kohn, S., Shin, T., Jackson, Y., Cardenas, M., McCann, R., Waterston, R. and Wilson, R.
The Washu-NCI, Mouse EST Project 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A1876493
uj59bl0.xl Sugano mouse liver mlia Mus musculus cDNA clone
IMAGE:1924219 3' similar to gb:x57025_rnal INSULIN-LIKE GROWTH
FACTOR IA PRECURSOR (HUMAN); gb:x04482 Mouse mRNA for
preproinsulin-like growth factor IB (MOUSE);, mRNA sequence.
A1876493.1 GI:5550542
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MGI:980511
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a quality sequence stop: 257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       quality sequence stop:
Location/Qualifiers
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314 286 1810
/note="Organ: liver; Vector: pME185-FL3; Site_1: DraIII (CACTGTGTG); Site_2: DraIII (CACCATGTG); 1st strand cDNA was primed with an oligo(dT) primer [ATGTGGCCTTTTTTTTTTTTTTTTT]; double-stranded cDNA was primed with an oligo(dT); double-stranded cDNA was ligated to a DraIII adaptor [TGTTGGCCTACTGG]; digested and cloned into distinct DraIII sites of the pME185-FL3 vector (5' site CACTGTGTG, 3' site CACCATGTG). XhoI should be used to isolate the CDNA insert. Size selection was performed to exclude fragments <1.5kb. Library constructed by Dr. Sumio Sugano (University of Tokyo Institute of Medical Science). Custom primers for sequencing: 5' end primer CTTCTGCTCTAAAAGCTGCG and 3' end primer CGACCTGCAGCTCGAGCACA."
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                                                                                                                                                                                                                                                                                                                                                                                                                                /strain="C57BL"
/db_xref="taxon:10090"
/clone="IMAGE:1924219"
                                                                                                                                                                                                                                                                                                                                                                                /clone_lib="Sugano mouse
/sex="female"
                                                                                                                                                                                                                                                                                                                           /lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                        /dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Mus musculus"
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                                                           6001 Executive Blvd. F
20892-9643, USA
Tel: 301 443 1706
Fax: 301 443 9890
Email: mEST@mail.nih.c
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UI-M-BH3-auy-g-11-0-UI.s1 NIH_BMAP_M_S4 Mus musculus
UI-M-BH3-auy-g-11-0-UI 3', mRNA sequence.
                                                                                                                        Contact: Chin, H
National Institute of Mental Health
National Institute of Mental Health
 The sequence contained an oligo-dT track that was present in the oligonucleotide that was used to prime the synthesis of first strand cDNA and therefore this may represent a bonafide poly A tail. The sequence tag present in the cDNA between the NotI site
                                                                                                                                                                                                                                                                                    Mus musculus
Eukaryota; Metazoa;
                                                                                                                                                                           Genome Res. 6 (9), 97044477
                                                                                                                                                                                                                                        1 (bases 1 to 499)
Bonaldo, M.F., Lenno
                                             Email: mEST@mail.nih.gov
The sequence contained an oligo-dT track that was
                                                                                                                                                                                                           discovery
                                                                                                                                                                                                                         Normalization and subtraction:
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Mammalia; Eutheria; Rodentia; Sciurognathi;
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78.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seq primer: M13 Forward POLYA=Yes.
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                           AGTCTCA----GAGAAGGAAAGGAACTACATTTGAAGAACACAAGTAGAGGGAGTGCAGGA 346
//dev_stage="7"-3" and days"
//dev_stage="7"-
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TAG_TISSUE=pineal-glands
TAG_SEQ=CAGAC"
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/clone_lib="NIH_BMAP_M_S4"
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6:791-806, 1996)
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Pred. No. 1.9e
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A1169770 468 bp
EST215669 Normalized rat liver,
RLIAT07 3' end, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      9712, Medical Center Drive, Tel: (301)-838-3529
Fax: (301)-838-0208
Email: nhlee@ttgr.org
Seq primer: M13-21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1998)
Other_ESTs: TC50779
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The Institute for Genomic Research
9712, Medical Center Drive, Rockville, MD 20850, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Lee, NH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lee, N.H., Glodek, A., Chandra, I., Mason, T.M., Kerlavage, A.R. and Adams, M.D.
Rat Genome Project: Generation of a Rat EST
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                                                                              CGCACCCCTCAAGCCTGCCAAGTCAGCTCGCTCTGTCCGTGCCCAGCGCCACACCCGACAT
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GCCCAAGACTCAGAAGTCCCAGCCCCTATCGACACACAAGAAAAGGAAGCTGCAAAGGAG
                             GCCCAAGACCCAGAAGTATCAGCCCCCATCTACCAACAAGAACACGAAGTCTCA---GAG
                                                              TGCTCCGCTGAAGCCTACAAAGTCAGCTCGTTCCATCCGGGCCCAGCGCCACACTGACAT
                                                                                                                             GGGCATTGTGGATGAGTGTTGCTCCCGGAGCTGTGATCTGAGGAGGTTGGAGATGTACTG
                                                                                                                                                              AGGCATCGTGGATGAGTGCTGCTTCCGGAGCTGTGATCTAAGGAGGCTGGAGATGTATTG
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                                                                                                                                                                                                                                                                                                                                          /clone_lib="Normalized rat liver, Bentc
/note="Organ: liver; Vector: pT7T3Pac;
Site_2: NotI"
1 115 c 119 g 149 t
                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Rattus sp."
/db_xref="ATCC (inhost):2027570"
/db_xref="taxon:10118"
/clone="RLIAT07"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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Pred. No. 4.5e-71;
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This read i
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Marra, M., Hiller, L., Kucaba, T., Martin, J., Beck, C., Wylie, T. Underwood, K., Steptoe, M., Theising, B., Allen, M., Bowers, Y., F., B., Swaller, T., Gibbons, M., Pape, D., Harvey, N., Schurk, R., F., E., Kohn, S., Shin, T., Jackson, Y., Cardenas, M., McCann, R., Waterston, R. and Wilson, R.

Waterston, R. and Wilson, R.

The WashU-NCI Mouse EST Project 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (1999)
Contact: Marra M/MashU-NCI Mouse EST Project 1999
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          preproinsulin-like growth factor AI604642
AI604642.1 GI:4613809
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seq primer: ~40RP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This read is a RESEQUENCE of a previously This read has been verified (found to hit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mus musculus
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314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         quality sequence stop:
Location/Qualifiers
                                                                                                                  /Clone_lib="Stratagene mouse diaphragm (#937303)"
/tissue_type="diaphragm"
/dev_stage="adult"
/lab_host="SOLR (kanamycin resistant)"
/l
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone="IMAGE:1001007"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Mus musculus"
/db_xref="taxon:10090"
   53.1%;
75.6%;
Pred. No. 1.6e-70;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              882 bp
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REFERENCE
AUTHORS
TITLE
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AI478804/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           384;
                                                                                         Emmert Buck, M.D., Ph.D.

cDNA Library Preparation: M. Bento Soares, Ph.D.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome

Clone distribution: NCI-CGAP clone distribution

found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.lnh.gov/bbrp/image/image.html

Insert Length: 849 Std Error: 0.00

Seq Primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                         Onpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A1478804 A1478804 A30 bp mRNA linear EST 14-APR-1999 tm52e04.x1 NCI_CGAP_Kidl1 Homo sapiens cDNA clone IMAGE:2161758 3' similar to gb:X57025_rnal INSULIN-LIKE GROWTH FACTOR IA PRECURSOR
                                                                                                                                                                                                                                                                                                                                                                                                    Mammmalia; Eutheria; Primates; Catarrhini; Hominic
1 (Dases 1 to 430)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ESI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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                                                    High quality sequence stop: 411.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (HUMAN);, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGACCGGAGACGCTCTGCGGGGCTGAGCTGGTGGATGCTCTTCAGTTCGTGTGTGGAGAC
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                                                                                                                                                                                                                                                                                                                                                                                  Gene Index
/organism="Homo sapiens
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BASE COUNT
ORIGIN
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Best Local Similarity
Matches 320; Conserv
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     130
                                                        482 AT-GGCATTTCCCCCAATGAAATACACAAGTAAACAT 517
                                                                                                             190 CGAGTTACCTGTTAAACTTTGGAACACCTACCAAAAAATAAGTTTGATAACATTTAAAAG 131
                                                                                                                                           424 C-AGTTACCTG-TAAACATTGGAATACCGGCCAAAAAATAAGTTTGATCACATTTCAAAG 481
                                                                                                                                                                                                                     250 TAGGAAGACCCTCCTGAGGAGTGAAGAGTGACATGCCACCGCAGGATCCTTTGCTCTGCA 191
                                                                                                                                                                                                                                                           365 TA-GAAGACCCTTCTGAGGAGTGAAGAAGGACAGGCCCACCGCAGGACCCTTTGCTCTGCA 423
                                                                                                                                                                                                                                                                                                                                                                                           305 AAGGAAGTACATTTGAAGAACACAAGTAGAGGGAGTGCAGGAAACAAGAACTACAGGATG 364
                                                                                                                                                                                                                                                                                                                                                                                                                                                370 CCAAGACCCAGAAGTATCAGCCCCCATCTACCAACAAGAACACGAAGTCTCAGAGAAGGA 311
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              185 CACCCCTCAAGCCTGCCAAGTCAGCTCGCTCTGTCCGTGCCCAGCGCCACCACCGACATGC 244
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note="Organ: kidney; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Plasmid DNA from the normalized library NOI_CGAP_Kid3 was prepared, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from a pool of 5,000 clones made from the same library (cloneIDs 132376-132391, 1456007-1456775, and 1500552-1502855). Subtraction by Bento Soares and M. Fatima Bonaldo. "
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93 c 10
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